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PCT/US03/18934

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| ggcccaattt                                       | cccacatttt | ggacaaaat  |            |            |            | 569 |
| <210> 16<br><211> 971<br><212> DNA<br><213> Homo | o sapien   |            |            |            |            |     |
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| atgaagacta                                       | ttctcagcaa | tcagactgtc | gacattccag | aaaatggtat | gagacttgat | 60  |
| gtcttttact                                       | tacatcttta | ctgcacgttc | caagcgttgt | gtggcctgac | gagtgtgttc | 120 |
| tctcttctag                                       | tcgacattac | tctgaaggga | cgcacagtta | tcgtgaaggg | ccccagagga | 180 |
| accctgcgga                                       | gggacttcaa | tcacatcaat | gtagaactca | gccttcttgg | aaagaaaaaa | 240 |
| aagaggctcc                                       | gggttgacaa | atggtggggt | aacagaaagg | aactggctac | cgttcggact | 300 |
| atttgtagtc                                       | atgtacagaa | catgatcaag | ggtgttacac | tgggcttccg | ttacaagatg | 360 |
| aggtctgtgt                                       | atgctcactt | ccccatcaac | gttgttatcc | aggagaatgg | gtctcttgtt | 420 |
| gaaatccgaa                                       | atttcttggg | tgaaaaatat | atccgcaggg | ttcggatgag | accaggtgtt | 480 |
| gcttgttcag                                       | tatctcaagc | ccagaaagat | gaattaatcc | ttgaaggaaa | tgacattgag | 540 |
| cttgtttcaa                                       | attcagcggc | tttgattcag | caagccacaa | cagttaaaaa | caaggatatc | 600 |
| aggaaatttt                                       | tggatggtat | ctatgtctct | gaaaaaggaa | ctgttcagca | ggctgatgaa | 660 |
| taagatctaa                                       | gagttacctg | gctacagaaa | gaagatgcca | gatgacactt | aagacctact | 720 |
| tgtgatattt                                       | aaatgatgca | ataaaagacc | tattgatttg | gaccttcttc | ttaaaaaaag | 780 |
| aaaaaaaga  | caaagaacaa | catagagcaa | aaacgagcaa | gcaaaaaaca | gaagaacaca | 840 |
| gccccgggcg                                       | attttattgt | tgggcgggcg | gcgcgaaacc | agggcctcag | tcaacggcca | 900 |
| ggttgccata                                       | ggggtgtccc | gcccctttt  | ttttccccga | gtgcgaacac | ccggcgcccc | 960 |
| aatgagggac                                       | a          |            |            |            |            | 971 |
|  |            |            |            |            |            |     |
| <210> 17 <211> 422 <212> DNA <213> Home          | o sapien   |            |            |            |            |     |
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|  | aaggagacat |            |            |            |            |     |
|  | catatttatg |            |            |            |            | 120 |
| aatgtcacca                                       | ctatctggag | atttcgacgt | gttttcctct | ctgaatctgt | tatgaacacg | 180 |
| ttaattaaat                                       | ggattgagta | ataaatatat | aaggggtttg | +++++aaaaa | aaaacaacaa | 240 |

| aaaaaaaaa  | aaaaaaccc    | ctggggcgta   | ccccggggca  | aaagggtggt   | ccacggggtg   | 300 |
|--|--------------|--------------|-------------|--------------|--------------|-----|
| agacttggtt                                       | ccccggcgca   | aaatccccc    | acactactaa  | gaacaagagg   | gccacggagg   | 360 |
| agcagcacgc                                       | acagatcaca   | gcagaccgac   | acagatagca  | acacagagac   | acacacgcat   | 420 |
| ag   |              |              |             |              |              | 422 |
|  |              |              |             |              |              |     |
| <210> 18<br><211> 584<br><212> DNA<br><213> Home | o sapien     |              |             |              |              |     |
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| aagaattcac                                       | tagtaatcgc   | catcgtggtg   | tgttcttgac  | teegetgete   | gccatgtctt   | 60  |
| ctcacaagac                                       | tttcaggatt   | aagcgattcc   | tggccaagaa  | acaaaagcaa   | aatcgtccca   | 120 |
| ttccccagtg                                       | gattcggatg   | aaaactggaa   | ataaaatcag  | gtacaactcc   | aaaaggagac   | 180 |
| attggagaag                                       | aaccaagctg   | ggtctataag   | gaattgcaca  | tgagatggca   | cacatattta   | 240 |
| tgctgtctga                                       | aggtcacgat   | catgttacca   | tatcaagctg  | aaaatgtcac   | cactatctgg   | 300 |
| agatttcgac                                       | gtgttttcct   | ctctgaatct   | gttatgaaca  | cgttggttgg   | ctggattcag   | 360 |
| taataaatat                                       | gtaaggcctt   | tctttttaaa   | aaaaaaaaa   | aaaaaaaaa    | aaaaaaaac    | 420 |
| ccctggggcg                                       | taccccgggg   | caaaagggtg   | gtccacgggg  | tgagacttgg   | ttccccggcg   | 480 |
| caaaatcccc                                       | ccacactact   | aagaacaaga   | gggccacgga  | ggagcagcac   | gcacagatca   | 540 |
| cagcagaccg                                       | acacagatag   | caacacagag   | acacacacgc  | atag         |              | 584 |
|  |              |              |             |              |              |     |
| <400> 19 acaatattga                              | acatttttct   | atatcctttg   | atatctgcaa  | gcctgatttt   | cagtagctgg   | 60  |
| aaatggaaag                                       | g gccaaattta | ttatctaatt   | ttatacatta  | ggacatgtgt   | ataatgtcca   | 120 |
| attttatact                                       | gttataagtc   | acactatgat   | gaacattttt  | gtacataact   | aaccatattt   | 180 |
| cagttcatt  | ctttaggtta   | ttatatatco   | acagatatga  | cattcaattc   | : tataaaaatt | 240 |
| atgtacatti                                       | taatttattt   | tatttttgta   | catgggaago  | tcctatctta   | actcattaaa   | 300 |
| ttcaataaa  | tttgtatttc   | : tacaacagaa | agccaacaaa  | a gggagttgtt | agtacatatt   | 360 |
| tccaggaat  | g aagttgtctg | gatgcagcta   | atgcctccat  | agaactgaca   | gtgctgaatt   | 420 |
| tacgaaatg  | g aaagagttct | ggaaaagcaa   | ı gaaaaaagt | cttgtttgaa   | a accccacgtc | 480 |
| tactgtagg  | c acagaaggga | atggaggcat   | ctgagcatt   | tattttccat   | : ctctacagca | 540 |
| cctcagaac  | a cctacattt  | attttttt     | ttctcagaaa  | a tgtcttaata | a agaggactgc | 600 |

| agtgtactca              | agtttcccaa   | tgacagggta                              | gggatgccaa | ccttctcttt | cattggcagc   | 660 |
|-------------------------|--------------|---|------------|------------|--------------|-----|
| tcatagtatc              | caagtttctc   | aaaaccctaa                              | gccatcttat | ttgttctttg | gaactttgtg   | 720 |
| gcctaccaca              | gtgcaatctc   | atcggtg                                 |            |            |              | 747 |
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| <210> 20<br><211> 766   |              |   |            |            |              |     |
| <212> DNA<br><213> Homo | sapien       |   |            |            |              |     |
| <400> 20                |              | )                                       |            |            | cagtaggtgg   | 60  |
|                         | acatttttct   |   |            |            |              |     |
|                         | gccaaattta   |   |            |            |              | 120 |
| attttatact              | gttataagtc   | acactatgat                              | gaacattttt | gtacataact | aaccatattt   | 180 |
| cagttcattt              | ctttaggtta   | ttatatatcc                              | acagatatga | cattcaattc | tataaaaatt   | 240 |
| atgtacattt              | taatttattt   | tatttttgta                              | catgggaagc | tcctatctta | actcattaaa   | 300 |
| ttcaataaat              | tttgtatttc   | tacaacagaa                              | agccaacaaa | gggagttgtt | agtacatatt   | 360 |
| tccaggaatg              | aagttgtctg   | gatgcagcta                              | atgcctccat | agaactgaca | gtgctgaatt   | 420 |
| tacgaaatgg              | aaagagttct   | ggaaaagcaa                              | gaaaaaaagt | cttgtttgaa | acccacgtc    | 480 |
| tactgtaggc              | acagaaggga   | atggaggcat                              | ctgagcattt | tattttccat | ctctacagca   | 540 |
| cctcagaaca              | cctacatttt   | atttttttc                               | ttctcagaaa | tgtcttaata | agaggactgc   | 600 |
| agtgtactca              | agtttcccaa   | tgacagggta                              | gggatgccaa | ccttctctt  | cattggcagc   | 660 |
| tcatagtatc              | caagtttctc   | aaaaccctaa                              | gccatcttat | ttgttctttg | gaactttgtg   | 720 |
| gcctaccaca              | gtgcaattct   | cattcggtgt                              | ttaataactc | gagccg     |              | 766 |
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| <210> 21<br><211> 647   |              |   |            |            |              |     |
| <212> DNA<br><213> Hom  | o sapien     |   |            |            |              |     |
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|                         | catgaataca   |   |            |            |              | 60  |
| acgtttgcag              | agcagtgggc   | acaatgttta                              | caatgtatgt | gtatgtcact | ttcggtacct   | 120 |
| gtgaatgcat              | ggggacgtgc   | tgaacccgaa                              | aaaaagtgcc | tttccataag | gactgcaata   | 180 |
| gagagggcaa              | tttaccctgg   | tggtacacgg                              | aacctagatt | cactcctgcc | atgccttgcc   | 240 |
| aatagtaago              | tgcagggtgg   | g aacaagaaat                            | cacttgctct | ggggggaagg | gaggggggaa   | 300 |
| tgggtgtgtc              | agctgggtag   | atacaaacco                              | tgaaaagaga | atccatgtgo | : tgctggcagg | 360 |
| anagatttt               | · taaaggtgtt | + | tcatatttco | gatttettt  | caggaaacat   | 420 |

|   |              | 20         |            |            |     |
|---|--------------|------------|------------|------------|-----|
| tcctgtggag ggaaaacgaa                                   | tatgaagata   | attttcagct | aattatctgg | gtgacccaga | 480 |
| atcgtgtata tggctatagg                                   | atagacttct   | taataatggc | aagtgacgtg | gccctgggga | 540 |
| aaggtgcttt atgtaccgtg                                   | tgtgcgtgta   | tgtgtgtgta | tctatacaag | tttgtcagct | 600 |
| ttggcatgac tgtttgtttg                                   | tctcgaaaac   | caataaactc | aaagttt    |            | 647 |
| <210> 22<br><211> 698<br><212> DNA<br><213> Homo sapien |              |            |            |            |     |
| <400> 22<br>actagcaccg ggcaagcaga                       | ı caacataatt | tatttccaga | aaacaacaga | atgaacatca | 60  |
| tcatgaatac atgaatcgg                                    |              |            |            |            | 120 |
| gagcagtggg cacaatgtt                                    |              |            |            |            | 180 |
| tggggacgtg ctgaacccg                                    |              |            |            |            | 240 |
|   |              |            |            |            | 300 |
| atttaccctg gtggtacac                                    |              |            |            |            |     |
| ctgcagggtg gaacaagaa                                    |              |            |            |            | 360 |
| cagctgggta gatacaaac                                    |              |            |            |            | 420 |
| ttaaagetet tteagaaae                                    | ctcatatttg   | gggtttcttt | tcaggaaaca | ttcctgtgga | 480 |
| gggaaaacga atatgaaga                                    | aattttcagc   | taattatctg | ggtgacccag | aatcgtgtat | 540 |
| atggctatag gatagactt                                    | c ttaataatgg | caagtgacgt | ggccctgggg | aaaggtgctt | 600 |
| tatgtaccgt gtgtgcgtg                                    | atgtgtgtgt   | atctatacaa | gtttgtcagc | tttggcatga | 660 |
| ctgtttgttt gtctcgaaa                                    | a ccaataaact | caaagttt   |            |            | 698 |
| <210> 23<br><211> 739<br><212> DNA<br><213> Homo sapien |              |            |            |            |     |
| <400> 23<br>taaacttaag gctaatgtt                        | agaagctttt   | gctaatgaga | ggaccatttg | ctaaatcggt | 60  |
| ataagtgcta cacatttgg                                    |              |            |            |            | 120 |
| gcagacaccg tctcctccc                                    |              |            |            |            | 180 |
| acactcacct ttctccttt                                    |              |            |            |            | 240 |
| aagctcctga tccagcagg                                    |              |            |            |            | 300 |
| cacccagcat ccaggcctc                                    |              |            |            |            | 360 |
|   |              |            |            |            | 420 |
| aaccetgcac cacatagac                                    |              |            |            |            |     |
| aatttgtgtc cttctgctt                                    | g gaactgtttc | ctttttagtt | tggtcaccct | cccagagctg | 480 |

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gtttcaatgg gggcataccc attatgggat gcagggcatc ctgcatcctg aggaattttt 540
tttcctccaa aaatgaaacc ttgaaatgag gacattgtcc tgtccacgga ctgcacaaca 600
acactgagcc tcaaggactc atactggcat ttttcttctt ttgcagagtg tgggcaccct 660
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tcataaagac agaatcaag 739

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<220>
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<sup>&</sup>lt;210> 25

<sup>&</sup>lt;211> 299

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Homo sapien

<sup>&</sup>lt;400> 25

28 60 ggcagcgcgg aggccgcacg atgcctggag ttactgtaaa agacgtgaac cagcaggagt 120 tegteagage tetggeagee tteeteaaaa agteegggga agetgaaagt eeeegaatgg 180 gtgggatacc gttcaagctg gccaaagcac aaaggagctt gctccctacg atgagaactg gttctacacg cggagctgct ttccaacagc ggcgggccac ctgttacctt ccgggggtgg 240 gcgctggggg ttgggcttcc attgaaccca aggattctat tgggggggaa cgttcagaa 299 <210> 26 <211> 1346 <212> DNA <213> Homo sapien <400> 26 60 ttttttttt ttgtgagcca gtgggaaaac caaggaggct aaaccataga gcctggagat gtgaaggaag tacaggtggg taagaaaggg agagccagat cacaagcacc ttgaaaccag 120 acactggttt ggggtcttca gcagtcctct gtcgaaatac atatattcag gggctgggtg 180 tggtggctca cacctgtaat cccagccctt tgggaggcag aggcaggcag attacttgag 240 gtcaagagtt caagacaagc ctggccaaca tggtaaaacc ccgtctctac caaaaatata 300 aaaaactagc cgggcgtggt ggcaggcacc tgattgtaat cccagctact cgggaggctg 360 aggcaggaga atcatttgaa cccagaaggc ggagattgca gtgagctgag atggcgccac 420 tgccactccc agcctgggcg acagagcaag agactcaaaa aagagaccca gaccaggatt 480 acgaatgagg caatttatta acccagcatg gtttgttcta atgcttcttg ttggcagctg 540 600 ccacctgtcc ggcgattctg tccagatctc tttgtccctg aggtgtcagt ttgcggccgc catcttggtc cttttccacc attttcagcc cctccagggc ttggaggacc cggcgggcca 660 cactettgga geeteggetg aagtggetgg geatgaegee gtttetetga egteeeceat 720 780 agatettggt catggageca accecagege cacceeggag gtacaggtge egegetgtgg aagcagctcg cgtgtagaac cagttctcat cgtagggagc aagctctttg tgcttggcca 840 gcttgacggt atccacccat tcggggactt tcagcttccc ggactttttg aggaaggctg 900 960 ccagagetet gaegaactee tgetggttea egtettttae agtaacteea ggeategtge ggcctccgcg ctgccagcca ggggaaaggg aacgacgggg tttcccgggc gcacaagtcg 1020 ggcgtagggt ctcgcgagag ttccgaaagc tcgcgagagc gagggtagac gctgaggctc 1080 1140 cgcctctctc agggcgaaag ttcgtccccg cctagagggg agggtgtcta gtgaggggtg gagaggtaaa ggggagggcc aaggggtcgc gcgtggaggc ctgggtttcc tcccgcgttt 1200 ccttctcccg gagtgtaata gagagagat agagagctcc tgttcggagc tgggggaact 1260 1320 tggcttcgtt tgcgtcgttc gtggctggaa ggaacagtgg tggagaatac tatgatggcg

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| aaagtac        | ggg gcaggatggg                              | tgggcc     |            |            |              | 1346 |
|----------------|---|------------|------------|------------|--------------|------|
|                | 27<br>136<br>DNA<br>Homo sapien             |            |            |            |              |      |
| <222>          | misc_feature<br>(75)(75)<br>n=a, c, g, or t | =          |            | ·          |              |      |
| <400>          | 27<br>Jetg egaagggage                       | caacaaata  | tatacacata | tacaatacat | aat aat aaaa | 60   |
|                |   |            |            |            |              |      |
|                | cca gtttnctgat                              | caagaggaat | ageceeerge | ceccagagea | ataaagtcag   | 120  |
| ctggctt        | tct cacctg                                  |            |            |            |              | 136  |
| <212>          | 28<br>426<br>DNA<br>Homo sapien             |            |            |            |              |      |
|                | 28<br>gcc atttcctctc                        | tccagaggac | ctttcctgcc | taggactcat | cattgtcccc   | 60   |
| tccctgg        | cat tttttacacc                              | tggagcagcc | agaggacgca | tgcatggctc | ttcggaagcc   | 120  |
| ttctcct        | gcc acggcatgca                              | cccacacatg | cgagcctccc | gggtactgtc | atcctgaatt   | 180  |
| ctgagac        | cat ccagcacttc                              | ctttagtttt | gccctggtgc | tgttgacttt | tgtttactga   | 240  |
| agagtgt        | gct ggaggcagga                              | caagggacat | ggaaggetge | aatttaagag | tctaaaaggt   | 300  |
| tttagaa        | tcc tgaaggaggt                              | ttaacaagct | gaattgaaga | ataatacctt | tctcaactgg   | 360  |
| agagaat        | tta catgattgca                              | ttattgttaa | aattaacatc | tcatctatta | aaagcatttg   | 420  |
| tagatt         |   |            |            |            |              | 426  |
| <211><br><212> | 29<br>264<br>DNA<br>Homo sapien             |            |            |            |              |      |
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|                | aca aagctgaatc                              |            |            |            |              | 120  |
|                | gtg acactacacc                              |            |            |            |              | 180  |
|                | tct cagttcacgg                              |            |            |            |              | 240  |
|                | taa agaagatgtg                              |            |            |            | -            | 264  |
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<211> 265
<212> DNA
<213> Homo sapien
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                                                                     120
accaageega tgtttgtgeg tggagaaaga tegtetttee teenteenea tgaeeegget
                                                                     180
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      741
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                                                                     120
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                                                                     180
tattgttcag ttttctgtat cttgcgcttg tttagccctg aaccaggagc aacagggtca
                                                                     240
gcttctggag gttggttgga acaatacggc aagtgctcga aatgacatcc agagaaatct
                                                                     300
aaactgctgt gggttccgca gtgttaaccc aaatgacacc tgtctggcta gctgtgttaa
                                                                     360
aagtgaccac tcgtgctcgc catgtgctcc aatcatagga gaatatgctg gagaggtttt
                                                                     420
gagatttgtt ggtggcattg gcctgttctt cagttttaca gagatcctgg gtgtttggct
                                                                     480
gacctacaga tacaggaacc agaaagaccc ccgcgcgaat cctagtgcat tcctttgatg
                                                                     540
agaaaacaag gaagatttcc tttcgtatta tgatcttgtt cactttctgt aattttctgt
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| taagctccat  | ttgccagttt    | aaggaaggaa | acactatctg | gaaaagtacc | ttattgatag | 660  |
|---|---------------|------------|------------|------------|------------|------|
| tggaattata  | tatttaccta    | gtttctctac | agttttcttc | cgtgcgaaaa | atattganac | 720  |
| tgggcctgaa  | ccggggcacg    | g          |            |            |            | 741  |
| <210> 32<br><211> 1844<br><212> DNA<br><213> Homo | l<br>o sapien |            |            |            |            |      |
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| agccccgccc  | tccctcctca    | cetgeteetg | gggaaactac | accaaggccg | cegetetgge | 120  |
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| ctcccggatc  | ccggtcggcg    | gagcgcattt | atttgcatat | ttctaccttt | gttccccgcc | 240  |
| tgggccaggc  | cccaaaggca    | aggacaaagc | agctgtcagg | gaacctccgc | cggagtcgaa | 300  |
| tttacgtgca  | gctgccggca    | accacaggtt | ccaagatggt | ttgcgggggc | ttcgcgtgtt | 360  |
| ccaagaactg  | cctgtgcgcc    | ctcaacctgc | tttacacctt | ggttagtctg | ctgctaattg | 420  |
| gaattgctgc  | gtggggcatt    | ggcttcgggc | tgatttccag | tctccgagtg | gtcggcgtgg | 480  |
| tcattgcagt  | gggcatcttc    | ttgttcctga | ttgctttagt | gggtctgatt | ggagctgtaa | 540  |
| aacatcatca  | ggtgttgcta    | ttcttttata | tgattattct | gttacttgta | tttattgttc | 600  |
| agttttctgt  | atcttgcgct    | tgtttagccc | tgaaccagga | gcaacagggt | cagcttctgg | 660  |
| aggttggttg  | gaacaatacg    | gcaagtgctc | gaaatgacat | ccagagaaat | ctaaactgct | 720  |
| gtgggttccg  | aagtgttaac    | ccaaatgaca | cctgtctggc | tagctgtgtt | aaaagtgacc | 780  |
| actcgtgctc  | gccatgtgct    | ccaatcatag | gagaatatgc | tggagaggtt | ttgagatttg | 840  |
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| gaatggtata                                      | ctaattggag   | ctaaagacgc | ttttcaccag              | ttgtttattg   | gttggccgtg | 1740 |
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| gaaggacgag                                      | acattccaat   | agttcacaga | gtaatcaaag              | ttcatgaaaa   | agataatgga | 180  |
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| teteccegeg                                      | teeggettgg   | atatcttcgg | ggacctgaag              | aagatgaaca   | agcgccagct | 180  |
| ctattaccag                                      | gttttaaact   | tcgccatgat | cgtgtcttct              | gcactcatga   | tatggaaagg | 240  |
| cttgatcgtg                                      | ctcacaggca   | gtgagagccc | catcgtggtg              | gtgctgagtg   | gcagtatgga | 300  |
| gccggccttt                                      | cacagaggag   | acctcctgtt | cctcacaaat              | ttccgggaag   | acccaatcag | 360  |
| agctggtgaa                                      | atagttgttt   | ttaaagttga | aggacgagac              | attccaatag   | ttcacagagt | 420  |
| aatcaaagtt                                      | . catgaaaaag | ataatggaga | catcaaattt              | ctgactaaag   | gagataataa | 480  |
| tgaagttgat                                      | gatagagget   | tgtacaaaga | aggccagaac              | : tggctggaaa | agaaggacgt | 540  |
| ggtgggaaga                                      | ı gcaagagggt | ttttaccata | tgttggtatg              | gtcaccataa   | taatgaatga | 600  |
|   |              |            | L _ L L L ~ ~ ~ ~ ~ ~ ~ |              | tactaaaacq | 660  |

33

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34

WO 03/106648 PCT/US03/18934

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| tcacctcggc | cccggacacc | aggccggccc | cgggctccac | cgcccccca  | gcccayggtg | 720  |
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| tcacctcggc | cccggacacc | aggccggccc | cgggctccac | cgcccccca  | gcccacggtg | 780  |
| teacetegge | cccggacacc | aggccggccc | cgggctccac | cgccccccca | gcccacggtg | 840  |
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| tcacctcggc | cccggacacc | aggccggccc | cgggctccac | cgcccccca  | gcccacggtg | 1920 |
| tcacctcggc | cccggacacc | aggccggccc | cgggctccac | cgcccccca  | gcccacggtg | 1980 |
| tcacctcggc | cccggacacc | aggccggccc | cgggctccac | cgcccccca  | gcccacggtg | 2040 |
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| acctctgcca | gggctaccac | aaccccagcc | agcaagagca | ctccattctc | aattcccagc | 3060 |
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| cagagageee | tgagatagcg | gggatcctga | actggactga | ataaaacgtg | gtctcccact | 4140 |
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PCT/US03/18934 WO 03/106648 37

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| tgctggtctg t                             | gttctggtt  | gcgctggcca   | ttgtctatct   | cattgccttg | gtgagtgcag | 1800 |
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| agcccgggat a                             | acctaccatc | ctatgagcga   | gtaccccacc   | taccacaccc | atgggcgcta | 1980 |
| tgtgcccct a                              | agcagtaccg | atcgtagccc   | ctatgagaag   | gtgagattgg | ccccacaggc | 2040 |
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| tttcaccacc                               | accatgacac | cgggcaccca   | gtctcctttc   | tteetgetge | tgctcctcac | 120  |
| agtgcttaca                               | gctaccacag | cccctacacc   | cgcaacagtt   | gttacaggtt | ctggtcatgc | 180  |
| aagctctacc                               | ccaggtggag | aaaaggagac   | ttcggctacc   | cagagaagtt | cagtgcccag | 240  |
| ctctactgag                               | aagaatgctg | tgagtatgac   | cagcagcgta   | ctctccagcc | acagccccgg | 300  |
| ttcaggctcc                               | tccaccactc | agggacagga   | tgtcactctg   | gccccggcca | cggaaccagc | 360  |
| ttcaggttca                               | gctgccacct | ggggacagga   | tgtcacctcg   | gtcccagtca | ccaggccagc | 420  |
| cctgggctcc                               | accaccccgc | cagcccacga   | tgtcacctca   | gccccggaca | acaagccagc | 480  |
| cccgggctcc                               | accgccccc  | cageceaegg   | tgtcacctcg   | gccccggaca | ccaggccggc | 540  |
| cccgggctcc                               | accgccccc  | cageceaegg   | tgtcacctcg   | gccccggaca | ccaggccggc | 600  |
| cccgggctcc                               | accgcgcccg | cagcccacgg   | tgtcacctcg   | gccccggaca | ccaggccggc | 660  |
| cccgggctcc                               | accgccccc  | cagcccatgg   | tgtcacctcg   | gccccggaca | acaggcccgc | 720  |
| cttggcgctc                               | caccgcccct | ccagtccaca   | . atgtcacctc | ggcctcaggc | tctgcatcag | 780  |
| gctcagcttc                               | tactctggtg | cacaacggca   | . cctctgccag | ggctaccaca | accccagcca | 840  |
| gcaagagcac                               | tccattctca | . attcccagco | accactetga   | tactcctacc | accettgeca | 900  |
| gccatagcac                               | caagactgat | gccagtagca   | ctcaccatag   | cacggtacct | cctctcacct | 960  |
| cctccaatca                               | cagcacttct | ccccagttgt   | . ctactggggt | ctctttcttt | ttcctgtctt | 1020 |
| ttcacatttc                               | aaacctccag | tttaattcct   | ctctggaaga   | tcccagcaco | gactactacc | 1080 |

| а | agagctgca  | gagagacatt | tctgaaatgt | ttttgcagat | ttataaacaa | gggggttttc | 1140 |
|---|------------|------------|------------|------------|------------|------------|------|
| t | gggcctctc  | caatattaag | ttcaggccag | gatctgtggt | ggtacaattg | actctggcct | 1200 |
| t | ccgagaagg  | taccatcaat | gtccacgacg | tggagacaca | gttcaatcag | tataaaacgg | 1260 |
| a | agcagcctc  | tcgatataac | ctgacgatct | cagacgtcag | cgtgagtgat | gtgccatttc | 1320 |
| c | tttctctgc  | ccagtctggg | gctggggtgc | caggctgggg | catcgcgctg | ctggtgctgg | 1380 |
| t | ctgtgttct  | ggttgcgctg | gccattgtct | atctcattgc | cttggctgtc | tgtcagtgcc | 1440 |
| 9 | gccgaaagaa | ctacgggcag | ctggacatct | ttccagcccg | ggatacctac | catcctatga | 1500 |
| ç | gcgagtaccc | cacctaccac | acccatgggc | gctatgtgcc | ccctagcagt | accgatcgta | 1560 |
| ç | gcccctatga | gaaggtttct | gcaggtaatg | gtggcagcag | cctctcttac | acaaacccag | 1620 |
| c | cagtggcagc | cacttctgcc | aacttgtagg | ggcacgtcgc | ccgctgagct | gagtggccag | 1680 |
| ( | ccagtgccat | tccactccac | tcaggttctt | cagggccaga | gcccctgcac | cctgtttggg | 1740 |
| ( | ctggtgagct | gggagttcag | gtgggctgct | cacagcctcc | ttcagaggcc | ccaccaattt | 1800 |
| ( | ctcggacact | tctcagtgtg | tggaagctca | tgtgggcccc | tgaggctcat | gcctgggaag | 1860 |
| 1 | tgttgtgggg | gctcccagga | ggactggccc | agagagccct | gagatagcgg | ggatcctgaa | 1920 |
| ( | ctggactgaa | taaaacgtgg | tctcccactg | cga        |            |            | 1953 |
|   |            |            |            |            |            |            |      |

<210> 40

<211> 1738

<212> DNA

<213> Homo sapien

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328

| acaatgtcac                             | ctcggcctca   | ggctctgcat | caggctcagc | ttctactctg | gtgcacaacg | 780  |
|--|--------------|------------|------------|------------|------------|------|
| gcacctctgc                             | cagggctacc   | acaaccccag | ccagcaagag | cactccattc | tcaattccca | 840  |
| gccaccactc                             | tgatactcct   | accacccttg | ccagccatag | caccaagact | gatgccagta | 900  |
| gcactcacca                             | tagcacggta   | cctcctctca | cctcctccaa | tcacagcact | tctccccagt | 960  |
| tgtctactgg                             | ggtctctttc   | tttttcctgt | cttttcacat | ttcaaacctc | cagtttaatt | 1020 |
| cctctctgga                             | agatcccagc   | accgactact | accaagagct | gcagagagac | atttctgaaa | 1080 |
| tgtttttgca                             | gatttataaa   | caagggggtt | ttctgggcct | ctccaatatt | aagttcaggc | 1140 |
| caggatctgt                             | ggtggtacaa   | ttgactctgg | ccttccgaga | aggtaccatc | aatgtccacg | 1200 |
| acgtggagac                             | acagttcaat   | cagtataaaa | cggaagcagc | ctctcgatat | aacctgacga | 1260 |
| tctcagacgt                             | cagcgtgagt   | gatgtgccat | ttcctttctc | tgcccagtct | ggggctgggg | 1320 |
| tgccaggctg                             | gggcatcgcg   | ctgctggtgc | tggtctgtgt | tctggttgcg | ctggccattg | 1380 |
| tctatctcat                             | tgccttggct   | gtctgtcagt | gccgccgaaa | gaactacggg | cagctggaca | 1440 |
| tctttccagc                             | ccgggatacc   | taccatccta | tgagcgagta | ccccacctac | cacacccatg | 1500 |
| ggcgctatgt                             | gccccctagc   | agtaccgatc | gtagccccta | tgagaaggtt | tctgcaggta | 1560 |
| atggtggcag                             | cagcctctct   | tacacaaacc | cagcagtggc | agccacttct | gccaacttgt | 1620 |
| aggggcacgt                             | cgcccgctga   | gctgagtggc | cagccagtgc | cattccactc | cactcagggc | 1680 |
| tetetgggee                             | agtectectg   | ggagccccca | ccacaacact | tcccaggcat | ggaattcc   | 1738 |
| <210> 41 <211> 328 <212> DNA <213> Hom |              |            | ·          |            |            |      |
| <400> 41<br>tcatctcgag                 | r cggcggcgca | gtgtgaggcg | gecegggete | accgcgcccg | cagcccacgg | 60   |
|  | gcccggaca    |            |            |            |            | 120  |
|  |              |            |            |            | cagcccatgg | 180  |
|  | gtgccycgga   |            |            |            |            | 240  |
|  |              |            |            |            | aagtatatcg | 300  |

atggatcata cgctgtttcc gtgtgtga

<220>

<sup>&</sup>lt;210> 42 <211> 1030 <212> DNA <213> Homo sapien

<221> misc\_feature <222> (574)..(574) <223> n=a, c, g, or t

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| <400> 42           |                 |            | tt at acassa | + aaaaa aaaa | 60   |
|--------------------|-----------------|------------|--------------|--------------|------|
| ccgctccacc tctcaag |                 |            |              |              |      |
| tttcaccacc accatga | acac cgggcaccca | gtctcctttc | ttcctgctgc   | tgctcctcac   | 120  |
| agtgcttaca gctacca | acag cccctacacc | cgcaacagtt | gttacaggtt   | ctggtcatgc   | 180  |
| aagctctacc ccaggt  | ggag aaaaggagac | ttcggctacc | cagagaagtt   | cagtgcccag   | 240  |
| ctctactgag aagaat  | gctg tgagtatgac | cagcagcgta | ctctccagcc   | acagccccgg   | 300  |
| ttcaggctcc tccacc  | actc agggacagga | tgtcactctg | gccccggcca   | cggaaccagc   | 360  |
| ttcaggttca gctgcc  | acct ggggacagga | tgtcacctcg | gtcccagtca   | ccaggccagc   | 420  |
| cctgggctcc accacc  | ccgc cagcccacga | tgtcacctca | gccccggaca   | acaagccagc   | 480  |
| cccgggctcc accgcc  | ccgc ggccgatctt | gtggctcggg | cttgggtacc   | gcgtgcgtgc   | 540  |
| ccggtcttca gctgct  | tcta gtaggtgctc | accntacgca | gttactaact   | tacgactgag   | 600  |
| cgctgtcgct ttgcac  | taga cgatcgtgaa | ctgggaacac | ctcatgtgct   | gtcatcacaa   | 660  |
| tttattcgct ttgcgg  | cgcg atccccctgt | tcgcaagagg | gtggaagagg   | ccactgtgtg   | 720  |
| taccccgcga acttag  | atcg tcggcggtgc | tagactagat | cacccctttg   | cgcagagact   | 780  |
| gagagtattg gggacc  | caga aaacagaagc | tgggggttca | ggagttttgc   | acgacaaaga   | 840  |
| actacgatag cagaag  | actt gatggtactg | gtgacccaag | gagaaatctg   | gggatttaga   | 900  |
| ggccacctga aagata  | cgaa gatacaaata | cagtctgaga | tgctggggac   | ccaggagaca   | 960  |
| gaggtggaca gcttct  | aggg taccagagtc | agaggctgag | ggggacagaa   | cgctaaaata   | 1020 |
| ttagggaccc         |                 |            |              |              | 1030 |

41

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<210> 43 <211> 1918 <212> DNA

<213> Homo sapien

<400> 43
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gcgggcgggc gggagtggg gggaccggta taaagcggta ggcgcctgtg cccgctccac 120
ctctcaagca gccagcgcct gcctgaatct gttctgccc ctcccaccc atttcaccac 180
caccatgaca ccgggcaccc agtctccttt cttcctgctg ctgctcctca cagtgcttac 240
agttgttaca ggttctggtc atgcaagctc taccccaggt ggagaaaagg agacttcggc 300
tacccagaga agttcagtgc ccagctctac tgagaagaat gctgtgagta tgaccagcag 360

| cgtactctcc | agccacagcc | ccggttcagg | ctcctccacc | actcagggac | aggatgtcac | 420  |
|------------|------------|------------|------------|------------|------------|------|
| tetggeeeeg | gccacggaac | cagcttcagg | ttcagctgcc | acctggggac | aggatgtcac | 480  |
| ctcggtccca | gtcaccaggc | cagccctggg | ctccaccacc | ccgccagccc | acgatgtcac | 540  |
| ctcagccccg | gacaacaagc | cagccccggg | ctccaccgcc | ccccagccc  | acggtgtcac | 600  |
| ctcggccccg | gacaccaggc | cggccccggg | ctccaccgcc | ccccagccc  | atggtgtcac | 660  |
| ctcggccccg | gacaacaggc | ccgccttggg | ctccaccgcc | cctccagtcc | acaatgtcac | 720  |
| ctcggcctca | ggctctgcat | caggctcagc | ttctactctg | gtgcacaacg | gcacctctgc | 780  |
| cagggctacc | acaaccccag | ccagcaagag | cactccattc | tcaattccca | gccaccactc | 840  |
| tgatactcct | accacccttg | ccagccatag | caccaagact | gatgccagta | gcactcacca | 900  |
| tagcacggta | cctcctctca | cctcctccaa | tcacagcact | tctccccagt | tgtctactgg | 960  |
| ggtctctttc | tttttcctgt | cttttcacat | ttcaaacctc | cagtttaatt | cctctctgga | 1020 |
| agatcccagc | accgactact | accaagagct | gcagagagac | atttctgaaa | tgtttttgca | 1080 |
| gatttataaa | caagggggtt | ttctgggcct | ctccaatatt | aagttcaggc | caggatctgt | 1140 |
| ggtggtacaa | ttgactctgg | ccttccgaga | aggtaccatc | aatgtccacg | acgtggagac | 1200 |
| acagttcaat | cagtataaaa | cggaagcagc | ctctcgatat | aacctgacga | tctcagacgt | 1260 |
| cagcgtgagt | gatgtgccat | ttcctttctc | tgcccagtct | ggggctgggg | tgccaggctg | 1320 |
| gggcatcgcg | ctgctggtgc | tggtctgtgt | tetggttgeg | ctggccattg | tctatctcat | 1380 |
| tgccttggct | gtctgtcagt | gccgccgaaa | gaactacggg | cagctggaca | tctttccagc | 1440 |
| ccgggatacc | taccatccta | tgagcgagta | cccacctac  | cacacccatg | ggcgctatgt | 1500 |
| gccccctagc | agtaccgatc | gtagccccta | tgagaaggtt | tetgeaggta | atggtggcag | 1560 |
| cagcctctct | tacacaaacc | cagcagtggc | agccacttct | gccaacttgt | aggggcacgt | 1620 |
| cgcccgctga | gctgagtggc | cagccagtgc | cattccactc | cactcaggtt | cttcagggcc | 1680 |
| agagcccctg | caccctgttt | gggctggtga | gctgggagtt | caggtgggct | gctcacagcc | 1740 |
| tccttcagag | gccccaccaa | tttctcggac | acttctcagt | gtgtggaagc | tcatgtgggc | 1800 |
| ccctgagggc | tcatgcctgg | gaagtgttgt | ggtgggggct | cccaggagga | ctggcccaga | 1860 |
| gagccctgag | atagcgggga | tcctgaactg | gactgaataa | aacgtggtct | cccactgc   | 1918 |

<sup>&</sup>lt;210> 44 <211> 1755 <212> DNA <213> Homo sapien

<sup>&</sup>lt;220>

<sup>&</sup>lt;221> misc\_feature <222> (1682)..(1682)

<223> n=a, c, g, or t

<220>

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<221> misc\_feature

<222> (1733)..(1733)

<223> n≈a, c, g, or t

<400> 44

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43

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| cagcctctct tacacaaacc               | cagcagtggc | agccacttct | gccaacttgt  | aggggcacgt | 1620 |
|-------------------------------------|------------|------------|-------------|------------|------|
| cgcccgctga gctgagtggc               | cagccagtgc | cattccactc | cactcaggtt  | cttcaggcag | 1680 |
| ancctgacct gttggctgta               | gctggagtca | gtggtgtaag | ctcttcaagg  | ggncagtcat | 1740 |
| cgatatgtaa cgttc                    |            |            |             |            | 1755 |
| .010- 45                            |            |            |             |            |      |
| <210> 45<br><211> 1530<br><212> DNA |            |            |             |            |      |
| <213> Homo sapien                   |            |            |             |            |      |
| <400> 45                            | aaaatttat  | anastatana | at ant ange | tataatataa | 60   |
| taggaggtag gggaggggc                |            |            |             |            |      |
| gcgggcgggc ggggagtggg               |            |            |             |            | 120  |
| ctctcaagca gccagcgcct               |            |            |             |            | 180  |
| caccatgaca ccgggcaccc               |            | _          | _           |            | 240  |
| agttgttaca ggttctggtc               | atgcaagctc | taccccaggt | ggagaaaagg  | agacttcggc | 300  |
| tacccagaga agttcagtgc               | ccagctctac | tgagaagaat | gcttttaatt  | cctctctgga | 360  |
| agateccage accgaetact               | accaagagct | gcagagagac | atttctgaaa  | tgtttttgca | 420  |
| gatttataaa caagggggtt               | ttctgggcct | ctccaatatt | aagttcaggc  | caggatctgt | 480  |
| ggtggtacaa ttgactctgg               | ccttccgaga | aggtaccatc | aatgtccacg  | acgtggagac | 540  |
| acagttcaat cagtataaaa               | cggaagcagc | ctctcgatat | aacctgacga  | tctcagacgt | 600  |
| cagcgtgagt gatgtgccat               | ttcctttctc | tgcccagtct | ggggctgggg  | tgccaggetg | 660  |
| gggcatcgcg ctgctggtgc               | tggtctgtgt | tctggttgcg | ctggccattg  | tctatctcat | 720  |
| tgccttggct gtctgtcagt               | gccgccgaaa | gaactacggg | cagctggaca  | tctttccagc | 780  |
| ccgggatacc taccatccta               | tgagcgagta | ccccacctac | cacacccatg  | ggcgctatgt | 840  |
| gccccctagc agtaccgatc               | gtagccccta | tgagaaggtt | tctgcaggta  | atggtggcag | 900  |
| cagcctctct tacacaaacc               | cagcagtggc | agccacttct | gccaacttgt  | aggggcacgt | 960  |
| cgcccgctga gctgagtggc               | cagccagtgc | cattccactc | cactcaggtt  | cttcagggcc | 1020 |
| agagcccctg caccctgttt               | gggctggtga | gctgggagtt | caggtgggct  | gctcacagcc | 1080 |
| tecttcagag gececaegae               | tatttcagga | agttcgaacc | ccacctgtac  | tecetegaet | 1140 |
| ccaacagcga cgatgtggac               | tctctgacag | acgaggagat | cctgtccaag  | taccagctgg | 1200 |
| gcatgctgca cttcagcact               | cagtacgacc | tgctgcacaa | ccacctcacc  | gtgcgcgtga | 1260 |
| tcgaggccag ggacctgcca               | cctcccatct | cccacgatgg | ctcgcgccag  | gacatggcgc | 1320 |
| actccaaccc ctacgtcaag               | atctgtctcc | tgccagacca | gaagaactca  | aagcagaccg | 1380 |

| gggtcaaacg caagacccag                                    | aagcccgtgt | ttgaggagcg | ctacaccttc | gagatcccct | 1440 |
|--|------------|------------|------------|------------|------|
| tcctggaggc ccagaggagg                                    | accetgetee | tgaccgtggt | ggattttgat | aagttctccc | 1500 |
| gccactgtgt cattgggaaa                                    | gtttctgtgg |            |            |            | 1530 |
| <210> 46<br><211> 563<br><212> DNA<br><213> Homo sapien  |            |            |            |            |      |
| <400> 46<br>ttttgctttt ttgcacccag                        | aggcaaaatg | ggtggagcac | tatgcccagg | ggagcccttc | 60   |
| ccgaggagtc ccaggggtga                                    | gcctctgtgc | ccctaatcat | ctcctaggaa | tggagggtag | 120  |
| accgagaaag gctggcatag                                    | ggggaggttt | cccaggtaga | agaagaagtg | tcagcagacc | 180  |
| aggtttctgc aggtaatggt                                    | ggcagcagcc | tctcttacac | aaacccagca | gtggcagcca | 240  |
| cttctgccaa cttgtagggg                                    | cacgtcgccc | gctgagctga | gtggccagcc | agtgccattc | 300  |
| cactccactc aggttcttca                                    | gggccagagc | ccctgcaccc | tgtttgggct | ggtgagctgg | 360  |
| gagttcaggt gggctgctca                                    | cagcctcctt | cagaggcccc | accaatttct | cggacacttc | 420  |
| tcagtgtgtg gaagctcatg                                    | tgggcccctg | agggctcatg | cctgggaagt | gttgtggtgg | 480  |
| gggctcccag gaggactggc                                    | ccagagagcc | ctgagatagc | ggggatcctg | aactggactg | 540  |
| aataaaacgt ggtctcccac                                    | tgc        | -          |            |            | 563  |
| <210> 47<br><211> 1945<br><212> DNA<br><213> Homo sapien |            |            |            |            |      |
| <400> 47<br>taggaggtag gggagggggc                        | ggggttttgt | cacctgtcac | ctgctccggc | tgtgctatgg | 60   |
| gcgggcgggc ggggagtggg                                    | gggaccggta | taaagcggta | ggcgcctgtg | cccgctccac | 120  |
| ctctcaagca gccagcgcct                                    | gcctgaatct | gttctgcccc | ctccccaccc | atttcaccac | 180  |
| caccatgaca ccgggcaccc                                    | agtctccttt | cttcctgctg | ctgctcctca | cagtgcttac | 240  |
| agctaccaca gcccctaaac                                    | ccgcaacagt | tgttacaggt | tctggtcatg | caagctctac | 300  |
| cccaggtgga gaaaaggaga                                    | cttcggctac | ccagagaagt | tcagtgccca | gctctactga | 360  |
| gaagaatget gtgagtatga                                    | ccagcagcgt | actctccagc | cacageceeg | gttcaggctc | 420  |
| ctccaccact cagggacagg                                    | atgtcactct | ggccccggcc | acggaaccag | cttcaggttc | 480  |
| agctgccacc tggggacagg                                    | atgtcacctc | ggtcccagtc | accaggccag | ccctgggctc | 540  |
| caccaccccg ccagcccacg                                    | atgtcacctc | agccccggac | aacaagccag | ccccgggctc | 600  |

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|  |            | 46         |            |            |      |
|--|------------|------------|------------|------------|------|
| caccgcccc ccagcccacg                                     | gtgtcacctc | ggccccggac | accaggccgg | ccccgggctc | 660  |
| caccgccccc ccagcccatg                                    | gtgtcacctc | ggccccggac | aacaggcccg | ccttgggctc | 720  |
| caccgcccct ccagtccaca                                    | atgtcacctc | ggcctcaggc | tctgcatcag | gctcagcttc | 780  |
| tactctggtg cacaacggca                                    | cctctgccag | ggctaccaca | accccagcca | gcaagagcac | 840  |
| tccattctca attcccagcc                                    | accactctga | tactcctacc | accettgeca | gccatagcac | 900  |
| caagactgat gccagtagca                                    | ctcaccatag | cacggtacct | cctctcacct | cctccaatca | 960  |
| cagcacttct ccccagttgt                                    | ctactggggt | ctctttcttt | ttcctgtctt | ttcacatttc | 1020 |
| aaacctccag tttaattcct                                    | ctctggaaga | tcccagcacc | gactactacc | aagagctgca | 1080 |
| gagagacatt tctgaaatgt                                    | ttttgcagat | ttataaacaa | gggggtttc  | tgggcctctc | 1140 |
| caatattaag ttcaggccag                                    | gatctgtggt | ggtacaattg | actctggcct | tccgagaagg | 1200 |
| taccatcaat gtccacgacg                                    | tggagacaca | gttcaatcag | tataaaacgg | aagcagcctc | 1260 |
| tcgatataac ctgacgatct                                    | cagacgtcag | cgtgagtgat | gtgccatttc | ctttctctgc | 1320 |
| ccagtctggg gctggggtgc                                    | caggctgggg | categegetg | ctggtgctgg | tctgtgttct | 1380 |
| ggttgcgctg gccattgtct                                    | atctcattgc | cttggctgtc | tgtcagtgcc | gccgaaagaa | 1440 |
| ctacgggcag ctggacatct                                    | ttccagcccg | ggatacctac | catcctatga | gcgagtaccc | 1500 |
| cacctaccac acccatgggc                                    | gctatgtgcc | ccctagcagt | accgatcgta | gcccctatga | 1560 |
| gaaggtttct gcaggtaatg                                    | gtggcagcag | cctctcttac | acaaacccag | cagtggcagc | 1620 |
| cacttctgcc aacttgtagg                                    | ggcacgtcgc | ccgctgagct | gagtggccag | ccagtgccat | 1680 |
| tccactccac tcaggttctt                                    | cagggccaga | gcccctgcac | cctgtttggg | ctggtgagct | 1740 |
| gggagttcag gtgggctgct                                    | cacagcctcc | ttcagaggcc | ccaccaattt | ctcggacact | 1800 |
| tctcagtgtg tggaagctca                                    | tgtgggcccc | tgagggctca | tgcctgggaa | gtgttgtggt | 1860 |
| gggggctccc aggaggactg                                    | gcccagagag | ccctgagata | gcggggatcc | tgaactggac | 1920 |
| tgaataaaac gtggtctccc                                    | actgc      |            |            |            | 1945 |
| <210> 48 <211> 1882 <212> DNA <213> Homo sapien <400> 48 |            |            | ·          |            |      |
| taggaggtag gggagggggc                                    | ggggttttgt | cacctgtcac | ctgctccggc | tgtgctatgg | 60   |
| gcgggcgggc ggggagtggg                                    | gggaccggta | taaagcggta | ggcgcctgtg | cccgctccac | 120  |
| ctctcaagca gccagcgcct                                    | gcctgaatct | gttctgcccc | ctccccaccc | atttcaccac | 180  |

caccatgaca ccgggcaccc agtctccttt cttcctgctg ctgctcctca cagtgcttac

|            |            |            | 4/         |            |            |      |
|------------|------------|------------|------------|------------|------------|------|
| aggtggagaa | aaggagactt | cggctaccca | gagaagttca | gtgcccagct | ctactgagaa | 300  |
| gaatgctgtg | agtatgacca | gcagcgtact | ctccagccac | agccccggtt | caggctcctc | 360  |
| caccactcag | ggacaggatg | tcactctggc | cccggccacg | gaaccagctt | caggttcagc | 420  |
| tgccacctgg | ggacaggatg | tcacctcggt | cccagtcacc | aggccagccc | tgggctccac | 480  |
| caccccgcca | gcccacgatg | tcacctcagc | cccggacaac | aagccagccc | cgggctccac | 540  |
| cgcccccca  | gcccacggtg | tcacctcggc | cccggacacc | aggccggccc | cgggctccac | 600  |
| cgcccccca  | gcccatggtg | tcacctcggc | cccggacaac | aggcccgcct | tgggctccac | 660  |
| cgcccctcca | gtccacaatg | tcacctcggc | ctcaggctct | gcatcaggct | cagcttctac | 720  |
| tctggtgcac | aacggcacct | ctgccagggc | taccacaacc | ccagccagca | agagcactcc | 780  |
| attctcaatt | cccagccacc | actctgatac | tcctaccacc | cttgccagcc | atagcaccaa | 840  |
| gactgatgcc | agtagcactc | accatagcac | ggtacctcct | ctcacctcct | ccaatcacag | 900  |
| cacttctccc | cagttgtcta | ctggggtctc | tttcttttc  | ctgtctttc  | acatttcaaa | 960  |
| cctccagttt | aattcctctc | tggaagatcc | cagcaccgac | tactaccaag | agctgcagag | 1020 |
| agacatttct | gaaatgtttt | tgcagattta | taaacaaggg | ggttttctgg | gcctctccaa | 1080 |
| tattaagttc | aggccaggat | ctgtggtggt | acaattgact | ctggccttcc | gagaaggtac | 1140 |
| catcaatgtc | cacgacgtgg | agacacagtt | caatcagtat | aaaacggaag | cagcctctcg | 1200 |
| atataacctg | acgatctcag | acgtcagcgt | gagtgatgtg | ccatttcctt | tctctgccca | 1260 |
| gtctggggct | ggggtgccag | gctggggcat | cgcgctgctg | gtgctggtct | gtgttctggt | 1320 |
| tgcgctggcc | attgtctatc | tcattgcctt | ggctgtctgt | cagtgccgcc | gaaagaacta | 1380 |
| cgggcagctg | gacatctttc | cagcccggga | tacctaccat | cctatgagcg | agtaccccac | 1440 |
| ctaccacacc | catgggcgct | atgtgccccc | tagcagtacc | gatcgtagcc | cctatgagaa | 1500 |
| ggtttctgca | ggtaatggtg | gcagcagcct | ctcttacaca | aacccagcag | tggcagccac | 1560 |
| ttctgccaac | ttgtaggggc | acgtcgcccg | ctgagctgag | tggccagcca | gtgccattcc | 1620 |
| actccactca | ggttcttcag | ggccagagcc | cctgcaccct | gtttgggctg | gtgagctggg | 1680 |
| agttcaggtg | ggctgctcac | agcctccttc | agaggcccca | ccaatttctc | ggacacttct | 1740 |
| cagtgtgtgg | aagctcatgt | gggcccctga | gggctcatgc | ctgggaagtg | ttgtggtggg | 1800 |
| ggctcccagg | aggactggcc | cagagagccc | tgagatagcg | gggatcctga | actggactga | 1860 |
| ataaaacgtg | gtctcccact | gc         |            |            |            | 1882 |
|            |            |            |            |            |            |      |

<sup>&</sup>lt;210> 49 <211> 1930 <212> DNA <213> Homo sapien

| <400> 49   |            |            |            |            |            |      |
|------------|------------|------------|------------|------------|------------|------|
| gtcgctctag | aggacccctc | ataggttcgc | agggccatga | gccaaggcct | atgggcagag | 60   |
| agaaggaggc | tgctgcaggg | aaggaggcgg | ccaacccagg | ggttactgag | gctgcccact | 120  |
| ccccagtcct | cctggtatta | tttctctggt | ggccagagct | tatattttct | tcttgctctt | 180  |
| atttttcctt | cataaagacc | caaccctatg | actttaactt | cttacagcta | ccacagcccc | 240  |
| taaacccgca | acagttgtta | caggttctgg | tcatgcaagc | tctaccccag | gtggagaaaa | 300  |
| ggagacttcg | gctacccaga | gaagttcagt | gcccagctct | actgagaaga | atgctgtgag | 360  |
| tatgaccagc | agcgtactct | ccagccacag | ccccggttca | ggctcctcca | ccactcaggg | 420  |
| acaggatgtc | actctggccc | cggccacgga | accagettea | ggttcagctg | ccacctgggg | 480  |
| acaggatgtc | acctcggtcc | cagtcaccag | gccagccctg | ggctccacca | ccccgccagc | 540  |
| ccacgatgtc | acctcagccc | cggacaacaa | gccagccccg | ggctccaccg | ccccccage  | 600  |
| ccacggtgtc | acctcggccc | cggacaccag | gccggccccg | ggctccaccg | ccccccagc  | 660  |
| ccatggtgtc | acctcggccc | cggacaacag | gcccgccttg | ggctccaccg | cccctccagt | 720  |
| ccacaatgtc | acctcggcct | caggctctgc | atcaggctca | gcttctactc | tggtgcacaa | 780  |
| cggcacctct | gccagggcta | ccacaacccc | agccagcaag | agcáctccat | tctcaattcc | 840  |
| cagccaccac | tctgatactc | ctaccaccct | tgccagccat | agcaccaaga | ctgatgccag | 900  |
| tagcactcac | catagcacgg | tacctcctct | cacctcctcc | aatcacagca | cttctcccca | 960  |
| gttgtctact | ggggtetett | tettttteet | gtcttttcac | atttcaaacc | tccagtttaa | 1020 |
| ttcctctctg | gaagatccca | gcaccgacta | ctaccaagag | ctgcagagag | acatttctga | 1080 |
| aatgttttg  | cagatttata | aacaaggggg | ttttctgggc | ctctccaata | ttaagttcag | 1140 |
| gccaggatct | gtggtggtac | aattgactct | ggccttccga | gaaggtacca | tcaatgtcca | 1200 |
| cgacgtggag | acacagttca | atcagtataa | aacggaagca | gcctctcgat | ataacctgac | 1260 |
| gatctcagac | gtcagcgtga | gtgatgtgcc | atttcctttc | tctgcccagt | ctggggctgg | 1320 |
| ggtgccaggc | tggggcatcg | cgctgctggt | gctggtctgt | gttctggttg | cgctggccat | 1380 |
| tgtctatctc | attgccttgg | ctgtctgtca | gtgccgccga | aagaactacg | ggcagctgga | 1440 |
| catctttcca | gcccgggata | cctaccatcc | tatgagcgag | taccccacct | accacaccca | 1500 |
| tgggcgctat | gtgcccccta | gcagtaccga | tcgtagcccc | tatgagaagg | tttctgcagg | 1560 |
| taatggtggc | agcagcctct | cttacacaaa | cccagcagtg | gcagccactt | ctgccaactt | 1620 |
| gtaggggcac | gtcgcccgct | gagctgagtg | gccagccagt | gccattccac | tccactcagg | 1680 |
| ttcttcaggg | ccagagcccc | tgcaccctgt | ttgggctggt | gagctgggag | ttcaggtggg | 1740 |
| ctgctcacag | cctccttcag | aggccccacc | aatttctcgg | acacttctca | gtgtgtggaa | 1800 |

| actcatataa  | gcccctgagg    | gctcatgcct    | gggaagtgtt   | gtggtgggg    | ctcccaggag   | 1860 |
|---|---------------|---------------|--------------|--------------|--------------|------|
|   |               |               | gatcctgaac   |              |              | 1920 |
|   | 949450005     | <b>~</b> 5~~~ | 5            |              |              | 1930 |
| ctcccactgc  |               |               |              |              |              |      |
| <210> 50<br><211> 1798<br><212> DNA<br><213> Home | 3<br>o sapien |               |              |              |              |      |
| <400> 50<br>taggaggtag                            | gggaggggc     | ggggttttgt    | cacctgtcac   | ctgctccggc   | tgtgctatgg   | 60   |
| gegggeggge  | ggggagtggg    | gggaccggta    | taaagcggta   | ggcgcctgtg   | cccgctccac   | 120  |
| ctctcaagca  | gccagcgcct    | gcctgaatct    | gttctgcccc   | ctccccaccc   | atttcaccac   | 180  |
| caccatgaca  | ccgggcaccc    | agtctccttt    | cttcctgctg   | ctgctcctca   | cagtgcttac   | 240  |
| agttgttaca  | ggttctggtc    | atgcaagctc    | taccccaggt   | ggagaaaagg   | agacttcggc   | 300  |
| tacccagaga  | agttcagtgc    | ccagctctac    | tgagaagaat   | gctgtgagta   | tgaccagcag   | 360  |
| cgtactctcc  | agccacagcc    | ccggttcagg    | ctcctccacc   | actcagggac   | aggatgtcac   | 420  |
| tetggeeeeg  | gccacggaac    | cagcttcagg    | ttcagctgcc   | acctggggac   | aggatgtcac   | 480  |
| ctcggtccca  | gtcaccaggc    | cagccctggg    | ctccaccacc   | ccgccagccc   | acgatgtcac   | 540  |
| ctcggccccg  | gacaacaggc    | ccgccttggg    | ctccaccgcc   | cctccagtcc   | acaatgtcac   | 600  |
| ctcggcctca  | ggctctgcat    | caggeteage    | ttctactctg   | gtgcacaacg   | gcacctctgc   | 660  |
| cagggctacc  | : acaaccccag  | ccagcaagag    | cactccattc   | tcaattccca   | gccaccactc   | 720  |
| tgatactcct  | accacccttg    | ccagccatag    | caccaagact   | gatgccagta   | gcactcacca   | 780  |
| tagcacggta  | cetectetea    | cctcctccaa    | tcacagcact   | tctccccagt   | tgtctactgg   | 840  |
| ggtctctttc  | tttttcctgt    | cttttcacat    | ttcaaacctc   | cagtttaatt   | cctctctgga   | 900  |
| agatcccago  | accgactact    | accaagagct    | gcagagagac   | atttctgaaa   | tgtttttgca   | 960  |
| gatttataaa  | a caagggggtt  | ttctgggcct    | ctccaatatt   | aagttcaggo   | caggatctgt   | 1020 |
| ggtggtacaa  | ttgactctgg    | ccttccgaga    | a aggtaccato | : aatgtccacg | g acgtggagac | 1080 |
| acagttcaat  | cagtataaa     | cggaagcag     | c ctctcgatat | aacctgacga   | tctcagacgt   | 1140 |
| cagcgtgagt  | gatgtgccat    | ttcctttct     | c tgcccagtct | ggggctgggg   | tgccaggctg   | 1200 |
| gggcatcgc   | g ctgctggtgd  | tggtctgtgt    | t tetggttgeg | g ctggccattg | g tctatctcat | 1260 |
| tgccttggc   | t gtctgtcagt  | ; gccgccgaa   | a gaactacggg | g cagctggace | a tctttccagc | 1320 |
| ccgggatac   | c taccatcct   | a tgagcgagt   | a ccccacctac | cacacccat    | g ggcgctatgt | 1380 |
| geceetag  | c agtaccgate  | gtagcccct     | a tgagaaggtt | tetgeaggta   | a atggtggcag | 1440 |

| cageetetet  | tacacaaacc    | cagcagtggc | agccacttct | gccaacttgt | aggggcacgt | 1500 |
|---|---------------|------------|------------|------------|------------|------|
| cgcccgctga  | gctgagtggc    | cagccagtgc | cattccactc | cactcaggtt | cttcagggcc | 1560 |
| agagcccctg  | caccctgttt    | gggctggtga | gctgggagtt | caggtgggct | gctcacagcc | 1620 |
| tccttcagag  | gccccaccaa    | tttctcggac | acttctcagt | gtgtggaagc | tcatgtgggc | 1680 |
| ccctgagggc  | tcatgcctgg    | gaagtgttgt | ggtgggggct | cccaggagga | ctggcccaga | 1740 |
| gagccctgag  | atagcgggga    | tectgaactg | gactgaataa | aacgtggtct | cccactgc   | 1798 |
| <210> 51<br><211> 1312<br><212> DNA<br><213> Homo | 2<br>o sapien |            |            |            |            |      |
|   | gggaggggc     | ggggttttgt | cacctgtcac | ctgctccggc | tgtgctatgg | 60   |
| gcgggcgggc  | ggggagtggg    | gggaccggta | taaagcggta | ggcgcctgtg | cccgctccac | 120  |
| ctctcaagca  | gccagcgcct    | gcctgaatct | gttctgcccc | ctccccaccc | atttcaccac | 180  |
| caccatgaca  | ccgggcaccc    | agtctccttt | cttcctgctg | ctgctcctca | cagtgcttac | 240  |
| agttgttaca  | ggttctggtc    | atgcaagctc | taccccaggt | ggagaaaagg | agacttcggc | 300  |
| tacccagaga  | agttcagtgc    | ccagctctac | tgagaagaat | gctttgtcta | ctggggtctc | 360  |
| tttcttttc   | ctgtctttc     | acatttcaaa | cctccagttt | aattcctctc | tggaagatcc | 420  |
| cagcacegac  | tactaccaag    | agctgcagag | agacatttct | gaaatgtttt | tgcagattta | 480  |
| taaacaaggg  | ggttttctgg    | gcctctccaa | tattaagttc | aggccaggat | ctgtggtggt | 540  |
| acaattgact  | ctggccttcc    | gagaaggtac | catcaatgtc | cacgacgtgg | agacacagtt | 600  |
| caatcagtat  | aaaacggaag    | cagcctctcg | atataacctg | acgatctcag | acgtcagcgt | 660  |
| gagtgatgtg  | ccatttcctt    | tctctgccca | gtctggggct | ggggtgccag | gctggggcat | 720  |
| cgcgctgctg  | gtgctggtct    | gtgttctggt | tgcgctggcc | attgtctatc | tcattgcctt | 780  |
| ggctgtctgt  | cagtgccgcc    | gaaagaacta | cgggcagctg | gacatctttc | cagcccggga | 840  |
| tacctaccat  | cctatgagcg    | agtaccccac | ctaccacacc | catgggcgct | atgtgccccc | 900  |
| tagcagtacc  | gatcgtagcc    | cctatgagaa | ggtttctgca | ggtaatggtg | gcagcagcct | 960  |
| ctcttacaca  | aacccagcag    | tggcagccac | ttctgccaac | ttgtaggggc | acgtcgcccg | 1020 |
| ctgagctgag  | tggccagcca    | gtgccattcc | actccactca | ggttcttcag | ggccagagcc | 1080 |
| cctgcaccct  | gtttgggctg    | gtgagctggg | agttcaggtg | ggctgctcac | agcctccttc | 1140 |
| agaggcccca  | ccaatttctc    | ggacacttct | cagtgtgtgg | aagctcatgt | gggcccctga | 1200 |
| gggctcatgc  | ctgggaagtg    | ttgtggtggg | ggctcccagg | aggactggcc | cagagagccc | 1260 |
|   |               |            |            |            |            |      |

| tgagatagcg  | gggatcctga   | actggactga   | ataaaacgtg | gtctcccact   | gc           | 1312 |
|---|--------------|--------------|------------|--------------|--------------|------|
| <210> 52<br><211> 2094<br><212> DNA<br><213> Homo | sapien       |              |            |              |              |      |
| <400> 52<br>taggaggtag                            | gggaggggc    | ggggttttgt   | cacctgtcac | ctgctccggc   | tgtgctatgg   | 60   |
| gcgggcgggc  | ggggagtggg   | gggaccggta   | taaagcggta | ggcgcctgtg   | cccgctccac   | 120  |
| ctctcaagca  | gccagcgcct   | gcctgaatct   | gttctgcccc | ctccccaccc   | atttcaccac   | 180  |
| caccatgaca  | ccgggcaccc   | agtctccttt   | cttcctgctg | ctgctcctca   | cagtgcttac   | 240  |
| agctaccaca  | gcccctaaac   | ccgcaacagt   | tgttacaggt | tctggtcatg   | caagctctac   | 300  |
| cccaggtgga  | gaaaaggaga   | cttcggctac   | ccagagaagt | tcagtgccca   | gctctactga   | 360  |
| gaagaatgct  | gtgagtatga   | ccagcagcgt   | actctccagc | cacagccccg   | gttcaggctc   | 420  |
| ctccaccact  | cagggacagg   | atgtcactct   | ggccccggcc | acggaaccag   | cttcaggttc   | 480  |
| agctgccacc  | tggggacagg   | atgtcacctc   | ggtcccagtc | accaggccag   | ccctgggctc   | 540  |
| caccaccccg  | ccagcccacg   | atgtcacctc   | agccccggac | aacaagccag   | ccccgggctc   | 600  |
| caccgccccc  | ccagcccacg   | gtgtcacctc   | ggccccggac | accaggccgg   | ccccgggctc   | 660  |
| caccgccccc  | ccagcccatg   | gtgtcacctc   | ggccccggac | aacaggcccg   | ccttgggctc   | 720  |
| caccgcccct  | ccagtccaca   | atgtcacctc   | ggcctcaggc | tctgcatcag   | gctcagcttc   | 780  |
| tactctggtg  | cacaacggca   | cctctgccag   | ggctaccaca | accccagcca   | gcaagagcac   | 840  |
| tccattctca  | attcccagcc   | accactctga   | tactcctacc | accettgeca   | gccatagcac   | 900  |
| caagactgat  | gccagtagca   | ctcaccatag   | cacggtacct | cctctcacct   | cctccaatca   | 960  |
| cagcacttct  | ccccagttgt   | ctactggggt   | ctctttcttt | tteetgtett   | ttcacatttc   | 1020 |
| aaacctccag  | tttaattcct   | ctctggaaga   | tcccagcacc | gactactaco   | aagagctgca   | 1080 |
| gagagacatt  | tctgaaatgt   | ttttgcagat   | ttataaacaa | gggggtttt    | tgggcctctc   | 1140 |
| caatattaag  | ttcaggtaca   | gttctgggtg   | tggacccagt | gtggtggttg   | gagggtgggt   | 1200 |
| ggtggtcatg  | accgtaggga   | gggactggtg   | cacttaaggt | tgggggaaga   | gtgctgagcc   | 1260 |
| agagctggga  | cccgtggctg   | aagtgcccat   | ttccctgtga | ccaggccagg   | g atctgtggtg | 1320 |
| gtacaattga  | ctctggcctt   | ccgagaaggt   | accatcaat  | g tecaegaegt | ggagacacag   | 1380 |
| ttcaatcagt  | . ataaaacgga | agcagcctct   | cgatataaco | tgacgatcto   | agacgtcagc   | 1440 |
| gtgagtgatg  | tgccatttcc   | : tttctctgcc | cagtctggg  | g ctggggtgc  | aggctggggc   | 1500 |
| atcgcgctgc  | : tggtgctggt | ctgtgttctg   | gttgcgctgg | g ccattgtcta | a tctcattgcc | 1560 |

| ttggctgtct g | tcagtgccg  | ccgaaagaac   | tacgggcagc | tggacatctt | tccagcccgg   | 1620 |
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| gatacctacc a | tcctatgag  | cgagtacccc   | acctaccaca | cccatgggcg | ctatgtgccc   | 1680 |
| cctagcagta c | cgatcgtag  | cccctatgag   | aaggtttctg | caggtaatgg | tggcagcagc   | 1740 |
| ctctcttaca c | aaacccagc  | agtggcagcc   | acttctgcca | acttgtaggg | gcacgtcgcc   | 1800 |
| cgctgagctg a | gtggccagc  | cagtgccatt   | ccactccact | caggttcttc | agggccagag   | 1860 |
| cccctgcacc c | tgtttgggc  | tggtgagctg   | ggagttcagg | tgggctgctc | acagcctcct   | 1920 |
| tcagaggccc ( | caccaatttc | tcggacactt   | ctcagtgtgt | ggaagctcat | gtgggcccct   | 1980 |
| gagggctcat g | gcctgggaag | tgttgtggtg   | ggggctccca | ggaggactgg | cccagagagc   | 2040 |
| cctgagatag ( | eggggateet | gaactggact   | gaataaaacg | tggtctccca | ctgc         | 2094 |
| <400> 53     | sapien     |              |            |            |              |      |
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| gcgggcgggc   |            |              |            |            |              | 120  |
| ctctcaagca   |            |              |            |            |              | 180  |
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| agctaccaca   | gcccctaaac | ccgcaacagt   | tgttacaggt | tctggtcatg | caagctctac   | 300  |
| cccaggtgga   | gaaaaggaga | cttcggctac   | ccagagaagt | tcagtgccca | gctctactga   | 360  |
| gaagaatgct   | gtgagtatga | ccagcagcgt   | actctccagc | cacageeeeg | gttcaggctc   | 420  |
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| agctgccacc   | tggggacagg | atgtcacctc   | ggtcccagtc | accaggccag | ccctgggctc   | 540  |
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| caccgccccc   | ccagcccacg | gtgtcacctc   | ggccccggac | accaggccgg | g ccccgggctc | 660  |
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| caccgcccct   | ccagtccaca | atgtcaccto   | ggcctcaggc | tctgcatcag | g gctcagcttc | 780  |
| tactctggtg   | cacaacggca | cctctgccag   | ggctaccaca | acccagcca  | a gcaagagcac | 840  |
| tccattctca   | attcccagco | accactctga   | tactcctacc | accettgee  | a gccatagcac | 900  |
| caagactgat   | gccagtagca | ctcaccatag   | cacggtacct | cctctcacc  | t cctccaatca | 960  |
| cagcacttct   | cccagttgt  | ctactggggt   | ctctttctt  | ttcctgtct  | t ttcacatttc | 1020 |
| aaacctccag   | tttaattcct | : ctctggaaga | tcccagcaco | gactactac  | c aagagctgca | 1080 |

| • | gagagacatt | tctgaaatgt | ttttgcagat | ttataaacaa | gggggttttc | tgggcctctc | 1140 |
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|   | taccatcaat | gtccacgacg | tggagacaca | gttcaatcag | tataaaacgg | aagcagcctc | 1260 |
|   | tcgatataac | ctgacgatct | cagacgtcag | cgtgagtgat | gtgccatttc | ctttctctgc | 1320 |
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|   | ggttgcgctg | gccattgtct | atctcattgc | cttggctgtc | tgtcagtgcc | gccgaaagaa | 1440 |
|   | ctacgggcag | ctggacatct | ttccagcccg | ggatacctac | catcctatga | gcgagtaccc | 1500 |
|   | cacctaccac | acccatgggc | gctatgtgcc | ccctagcagt | accgatcgta | gcccctatga | 1560 |
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|   | gggagttcag | gtgggctgct | cacagcctcc | ttcagaggcc | ccaccaattt | ctcggacact | 1800 |
|   | tctcagtgtg | tggaagctca | tgtgggcccc | tgagggctca | tgcctgggaa | gtgttgtggt | 1860 |
|   | gggggctccc | aggaggactg | gcccagagag | ccctgagata | geggggatee | tgaactggac | 1920 |
|   | tgaataaaac | gtggtctccc | actgcaaaag | acataaaaaa | agaaaaagac | aaagacgagc | 1980 |
|   | aaaaagacaa | aaagaggcaa | aaacaacaaa | acacaacaaa | caaaaaaaag | cacacacaaa | 2040 |
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<211> 2194

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<213> Homo sapien

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PCT/US03/18934 WO 03/106648

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| caccgccccc ccagcccatg gtgtcacctc ggccccggac aacaggcccg ccttgggctc | 720  |
| caccgcccct ccagtccaca atgtcacctc ggcctcaggc tctgcatcag gctcagcttc | 780  |
| tactctggtg cacaacggca cctctgccag ggctaccaca accccagcca gcaagagcac | 840  |
| tccattctca attcccagcc accactctga tactcctacc acccttgcca gccatagcac | 900  |
| caagactgat gccagtagca ctcaccatag cacggtacct cctctcacct cctccaatca | 960  |
| cagcacttct ccccagttgt ctactggggt ctctttcttt ttcctgtctt ttcacatttc | 1020 |
| aaacctccag tttaattcct ctctggaaga tcccagcacc gactactacc aagagctgca | 1080 |
| gagagacatt tetgaaatgg tgagtategg cettteette eecatgetee eetgaageag | 1140 |
| ccatcagaac tgtccacacc ctttgcatca agcctgagtc ctttccctct caccccagtt | 1200 |
| ttttgcagat ttataaacaa gggggttttc tgggcctctc caatattaag ttcaggtaca | 1260 |
| gttctgggtg tggacccagt gtggtggttg gagggtgggt ggtggtcatg accgtaggga | 1320 |
| gggactggtg cacttaaggt tgggggaaga gtgctgagcc agagctggga cccgtggctg | 1380 |
| aagtgcccat ttccctgtga ccaggccagg atctgtggtg gtacaattga ctctggcctt | 1440 |
| ccgagaaggt accatcaatg tccacgacgt ggagacacag ttcaatcagt ataaaacgga | 1500 |
| agcagcetet egatataace tgaegatete agaegteage gtgagtgatg tgecatttee | 1560 |
| tttetetgee cagtetgggg etggggtgee aggetgggge ategegetge tggtgetggt | 1620 |
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| cccctatgag aaggtttctg caggtaatgg tggcagcagc ctctcttaca caaacccagc | 1860 |
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| cagtgccatt ccactccact caggttcttc agggccagag cccctgcacc ctgtttgggc | 1980 |
| tggtgagetg ggagtteagg tgggetgete acageeteet teagaggeee caceaattte | 2040 |
| teggacaett eteagtgtgt ggaageteat gtgggeeeet gagggeteat geetgggaag | 2100 |
| tgttgtggtg ggggctccca ggaggactgg cccagagagc cctgagatag cggggatcct | 2160 |
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| ctctcaagca             | gccagcgcct   | gcctgaatct   | gttctgcccc | ctccccaccc   | atttcaccac   | 180  |
| caccatgaca             | ccgggcaccc   | agtctccttt   | cttcctgctg | ctgctcctca   | cagtgcttac   | 240  |
| agttgttaca             | ggttctggtc   | atgcaagctc   | taccccaggt | ggagaaaagg   | agacttcggc   | 300  |
| tacccagaga             | agttcagtgc   | ccagctctac   | tgagaagaat | gctatcccag   | caccgactac   | 360  |
| taccaagagc             | tgcagagaga   | catttctgaa   | atggccagga | tctgtggtgg   | tacaattgac   | 420  |
| tctggccttc             | cgagaaggta   | ccatcaatgt   | ccacgacgtg | gagacacagt   | tcaatcagta   | 480  |
| taaaacggaa             | gcagcctctc   | gatataacct   | gacgatctca | gacgtcagcg   | tgagtgatgt   | 540  |
| gccatttcct             | ttctctgccc   | agtctggggc   | tggggtgcca | ggctggggca   | tegegetget   | 600  |
| ggtgctggtc             | tgtgttctgg   | ttgcgctggc   | cattgtctat | ctcattgcct   | tggctgtctg   | 660  |
| tcagtgccgc             | cgaaagaact   | acgggcagct   | ggacatcttt | ccagcccggg   | atacctacca   | 720  |
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| cgatcgtagc             | ccctatgaga   | aggtttctgc   | aggtaatggt | ggcagcagcc   | tctcttacac   | 840  |
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| gtggccagcc             | agtgccattc   | cactccactc   | aggttcttca | gggccagagc   | ccctgcaccc   | 960  |
| tgtttgggct             | ggtgagctgg   | gagttcaggt   | gggctgctca | cagcctcctt   | cagaggcccc   | 1020 |
| accaatttct             | . cggacacttc | tcagtgtgtg   | gaagctcatg | tgggcccctg   | agggctcatg   | 1080 |
| cctgggaagt             | gttgtggtgg   | gggctcccag   | gaggactggc | ccagagagcc   | ctgagatagc   | 1140 |
| ggggatcctg             | g aactggactg | aataaaacgt   | ggtctcccac | tgc          |              | 1183 |
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| ctctcaagc              | a gccagcgcct | gcctgaatct   | gttctgccc  | : ctccccaccc | atttcaccac   | 180  |
| caccatgaca             | a ccgggcacco | agtctccttt   | cttcctgctg | , ctgctcctca | cagtgcttac   | 240  |
| agctaccaca             | a gcccctaaac | c ccgcaacagt | tgttacaggt | : tctggtcatg | g caagctctac | 300  |
| cccaggtgg              | a gaaaaggaga | cttcggctac   | ccagagaagt | : tcagtgccca | gctctactga   | 360  |
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| ctccaccact | cagggacagg | atgtcactct   | ggccccggcc | acggaaccag   | cttcaggttc   | 480  |
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| cacegeceet | ccagtccaca | atgtcacctc   | ggcctcaggc | tctgcatcag   | gctcagcttc   | 780  |
| tactctggtg | cacaacggca | cctctgccag   | ggctaccaca | accccagcca   | gcaagagcac   | 840  |
| tccattctca | attcccagcc | accactctga   | tactcctacc | accettgeca   | gccatagcac   | 900  |
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| aaacctccag | tttaattcct | ctctggaaga   | tcccagcacc | gactactacc   | aagagctgca   | 1080 |
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| caatattaag | ttcaggccag | gatctgtggt   | ggtacaattg | actctggcct   | tccgagaagg   | 1200 |
| taccatcaat | gtccacgacg | tggagacaca   | gttcaatcag | tataaaacgg   | aagcagcctc   | 1260 |
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| taccgtgcta | tggtgagtgc | tactggcatc   | agtcttggtg | ctatggctgg   | caagggtggt   | 1380 |
|            |            | tgggaattga   |            |              |              | 1440 |
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| cactcaccat | agcacggtac | ctcctctcac   | ctcctccaat | cacagcactt   | ctccccagtt   | 1560 |
|            |            | tttcctgtc    |            |              |              | 1620 |
| tctctggaag | atcccagcac | : cgactactac | caagagctgc | : agagagacat | ttctgaaatg   | 1680 |
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|            |            |              |            |              | ggcagcagcc   | 1980 |
|            |            |              |            |              | cacgtcgccc   | 2040 |
|            |            |              |            |              | gggccagagc   | 2100 |
|            |            |              |            |              | cagcctcctt   | 2160 |
|            |            |              |            |              | g tgggcccctg | 2220 |
|            |            |              |            |              |              |      |

57

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58

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|----------------------------------|------------------------------------|------|
| ggacacttct cagtgtgtgg aagctcatgt | gggcccctga gggctcatgc ctgggaagtg   | 1620 |
| ttgtggtggg ggctcccagg aggactggcc | c cagagagece tgagatageg gggateetga | 1680 |
| actggactga ataaaacgtg gtctcccact | gc :                               | 1712 |

<210> 58 <211> 1605 <212> DNA <213> Homo sapien

4005 ER

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| ( | ctctcaagca | gccagcgcct   | gcctgaatct | gttctgcccc | ctccccaccc | atttcaccac | 180  |
| • | caccatgaca | ccgggcaccc   | agtctccttt | cttcctgctg | ctgctcctca | cagtgcttac | 240  |
| ä | agttgttaca | ggttctggtc   | atgcaagctc | taccccaggt | ggagaaaagg | agacttcggc | 300  |
| 1 | tacccagaga | agttcagtgc   | ccagctctac | tgagaagaat | gctgtgagta | tgaccagcag | 360  |
| , | cgtactctcc | agccacagcc   | ccggttcagg | ctcctccacc | actcagggac | aggatgtcac | 420  |
| 1 | tetggeeeeg | gccacggaac   | cagcttcagg | ttcagctgcc | acctggggac | aggatgtcac | 480  |
| , | ctcggtccca | gtcaccaggc   | cagccctggg | ctccaccacc | ccgccagccc | acgatgtcac | 540  |
|   | ctcggccccg | gacaacaggc   | ccgccttggg | ctccaccgcc | cctccagtcc | acaatgtcac | 600  |
|   | ctcggcctca | ggctctgcat   | caggctcagc | ttctactctg | gtgcacaacg | gcacctctgc | 660  |
|   | cagggctacc | acaaccccag   | ccagcaagag | cactccattc | tcaattccca | gccaccactc | 720  |
|   | tgatactcct | accacccttg   | ccagccatag | caccaagact | gatgccagta | gcactcacca | 780  |
|   | tagcacggta | cctcctctca   | cctcctccaa | tcacagcact | tctccccagt | tgtctactgg | 840  |
|   | ggtctctttc | ttttcctgt    | cttttcacat | ttcaaacctc | cagtttaatt | cctctctgga | 900  |
|   | agatcccago | accgactact   | accaagagct | gcagagagac | atttctgaaa | tgtgagtgat | 960  |
|   | gtgccatttc | ctttctctgc   | ccagtctggg | gctggggtgc | caggctgggg | categegetg | 1020 |
|   | ctggtgctgg | tctgtgttct   | ggttgcgctg | gccattgtct | atctcattgc | cttggctgtc | 1080 |
|   | tgtcagtgcc | gccgaaagaa   | ctacgggcag | ctggacatct | ttccagcccg | ggatacctac | 1140 |
|   | catcctatga | gcgagtaccc   | cacctaccac | acccatgggc | gctatgtgcc | ccctagcagt | 1200 |
|   | accgatcgta | gcccctatga   | gaaggtttct | gcaggtaatg | gtggcagcag | cctctcttac | 1260 |
|   | acaaacccag | r cagtggcagc | cacttctgcc | aacttgtagg | ggcacgtcgc | ccgctgagct | 1320 |
|   | gagtggccag | ccagtgccat   | tccactccac | tcaggttctt | cagggccaga | gcccctgcac | 1380 |

| 59  | -    |
|---|------|
| cctgtttggg ctggtgagct gggagttcag gtgggctgct cacagcctcc ttcagaggcc         | 1440 |
| ccaccaattt ctcggacact tctcagtgtg tggaagetca tgtgggeece tgagggetca         | 1500 |
| tgcctgggaa gtgttgtggt gggggctccc aggaggactg gcccagagag ccctgagata         | 1560 |
| gcggggatcc tgaactggac tgaataaaac gtggtctccc actgc                         | 1605 |
| <210> 59 <211> 1874 <212> DNA <213> Homo sapien                           |      |
| <400> 59 taggaggtag gggaggggc ggggttttgt cacctgtcac ctgctccggc tgtgctatgg | 60   |
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| ctctcaagca gccagcgcct gcctgaatct gttctgcccc ctccccaccc atttcaccac         | 180  |
| caccatgaca cegggeacee agteteettt etteetgetg etgeteetea eagtgettae         | 240  |
| agctaccaca gcccctaaac ccgcaacagt tgttacaggt tctggtcatg caagctctac         | 300  |
| cccaggtgga gaaaaggaga cttcggctac ccagagaagt tcagtgccca gctctactga         | 360  |
| gaagaatget gtgagtatga ceageagegt actetecage cacageeceg gttcaggete         | 420  |
| ctccaccact cagggacagg atgtcactct ggccccggcc acggaaccag cttcaggttc         | 480  |
| agetgecace tggggacagg atgteacete ggteecagte accaggecag ceetgggete         | 540  |
| caccacccg ccagcccacg atgtcacctc agccccggac aacaagccag ccccgggctc          | 600  |
| cacegecece ecageceacg gtgteacete ggeeceggae accaggeegg eccegggete         | 660  |
| caccgcccc ccagcccatg gtgtcacctc ggccccggac aacaggcccg ccttgggctc          | 720  |
| caccgcccct ccagtccaca atgtcacctc ggcctcaggc tctgcatcag gctcagcttc         | 780  |
| tactetggtg cacaacggca cetetgecag ggetaccaca accecageca geaagageac         | 840  |
| tocattotca attoccagoo accaetetga tacteetace accettgeca gecatageae         | 900  |
| caagactgat gccagtagca ctcaccatag cacggtacct cctctcacct cctccaatca         | 960  |

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gagagacatt tctgaaatgt ttttgcagat ttataaacaa gggggttttc tgggcctctc

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ccagtctggg gctggggtgc caggctgggg catcgcgctg ctggtgctgg tctgtgttct

1020

1080

1140

1200

1260

1320

60

| ggttgcgctg | gccattgtct | atctcattgc | cttggctgtc | tgtcagtgcc | gccgaaagaa | 1440 |
|------------|------------|------------|------------|------------|------------|------|
| ctacgggcag | ctggacatct | ttccagcccg | ggatacctac | catcctatga | gcgagtaccc | 1500 |
| cacctaccac | acccatgggc | gctatgtgcc | ccctagcagt | accgatcgta | gcccctatga | 1560 |
| gaaggtgaga | ttgggcccca | caggccaggg | gaagcagagg | gtttggctgg | gcaaggattc | 1620 |
| tgaagggggt | acttggaaaa | cccaaagagc | ttggaagagg | tgagaagtgg | cgtgaagtga | 1680 |
| gcaggggagg | gcctggcaag | gatgaggggc | agaggtcaga | ggagttttgg | gggacaggcc | 1740 |
| tgggaggaga | ctatggaaga | aaggggccct | caagagggag | tggccccact | gccagaattc | 1800 |
| ctaaaagatc | attggccgtc | cacattcatg | ctggctggcg | ctggctgaac | tggtgccacc | 1860 |
| gtggcagttt | tgtt       |            |            |            |            | 1874 |

<210> 60

<211> 1634

<212> DNA

<213> Homo sapien

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61

gagagacatt tetgaaatgt ttttgcagat ttataaacaa gggggtttte tgggeetete 1140 caatattaag ttcaggccag gatctgtggt ggtacaattg actctggcct tccgagaagg 1200 taccatcaat gtccacgacg tggagacaca gttcaatcag tataaaacgg aagcagcctc 1260 tcgatataac ctgacgatct cagacgtcag cgtgagtgat gtgccatttc ctttctctgc 1320 ccagtctggg gctggggtgc caggctgggg catcgcgctg ctggtgctgg tctgtgttct 1380 ggttgcgctg gccattgtct atctcattgc cttggctgtc tgtcagtgcc gccgaaagaa 1440 ctacgggcag ctggacatct ttccagcccg ggatacctac catcctatga gcgagtggag 1500 ggtgtagaag agaagaagaa ggaggttcct gctgtgccag aaacccttaa gaaaaagcga 1560 aggaatttcg cagagctgaa gatcaagcgc ctgagaaaga agttksccaa aagatgcttc 1620 1634 gaaaggcaag gagg

<210> 61

<211> 943

<212> DNA

<213> Homo sapien

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62

<211> 997
<212> DNA
<213> Homo sapien
<400> 62
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ctctcaagca gccagcgcct gcctgaatct gttctgcccc ctccccaccc atttcaccac 180

caccatgaca cegggcacec agteteettt etteetgetg etgeteetea cagtgettac 240

agttgttaca ggttctggtc atgcaagctc taccccaggt ggagaaaagg agacttcggc 300

tacccagaga agttcagtgc ccagctctac tgagaagaat getttgtcta etggggtetc 360

tttettttte etgtetttte acattteaaa eeteeagttt aatteetete tggaagatee 420

cagcaccgac tactaccaag agctgcagag agacatttct gaaatggctg tctgtcagtg 480

ccgccgaaag aactacgggc agctggacat ctttccagcc cgggatacct accatcctat 540

gagegagtac eccacetace acacecatgg gegetatgtg ecceetagea gtacegateg 600

tagcccctat gagaaggttt ctgcaggtaa tggtggcagc agcctctctt acacaaaccc 660

agcagtggca gccacttetg ccaacttgta ggggcacgtc gcccgctgag ctgagtggcc 720

agccagtgcc attccactcc actcaggttc ttcagggcca gagcccctgc accctgtttg 780

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cctqaactgg actgaataaa acgtggtctc ccactgc 997

<210> 63

<211> 548

<212> DNA

<213> Homo sapien

<400> 63

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<210> 64 <211> 1378 <212> DNA

<213> Homo sapien

<400> 64

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<sup>&</sup>lt;210> 65

<sup>&</sup>lt;211> 162

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| <212> DNA<br><213> Homo sapien                           |            |            |            |      |
|--|------------|------------|------------|------|
| <400> 65<br>gcggccgcct actactacta ctgctcgaat             | tcaagcttct | aacgatgtac | gggctcatgc | 60   |
| ctgggaagtg ttgtggtggg ggctcccagg                         | aggactggcc | cagagagccc | tgagatagcg | 120  |
| gggatcctga actggactga ataaaacgtg                         | gtctcccact | gc         |            | 162  |
| <210> 66<br><211> 1285<br><212> DNA<br><213> Homo sapien |            |            |            |      |
| <400> 66<br>taggaggtag gggagggggc ggggttttgt             | cacctgtcac | ctgctccggc | tgtgctatgg | 60   |
| gcgggcgggc ggggagtggg gggaccggta                         | taaagcggta | ggcgcctgtg | cccgctccac | 120  |
| ctctcaagca gccagcgcct gcctgaatct                         | gttctgcccc | ctccccaccc | atttcaccac | 180  |
| caccatgaca cogggcaccc agtctccttt                         | cttcctgctg | ctgctcctca | cagtgcttac | 240  |
| agctaccaca gcccctaaac ccgcaacagt                         | tgttacaggt | tctggtcatg | caagctctac | 300  |
| cccaggtgga gaaaaggaga cttcggctac                         | ccagagaagt | tcagtgccca | gctctactga | 360  |
| gaagaatget tttaatteet etetggaaga                         | tcccagcacc | gactactacc | aagagctgca | 420  |
| gagagacatt tctgaaatgt ttttgcagat                         | ttataaacaa | gggggtttc  | tgggcctctc | 480  |
| caatattaag ttcaggccag gatctgtggt                         | ggtacaattg | actctggcct | tccgagaagg | 540  |
| taccatcaat gtccacgacg tggagacaca                         | gttcaatcag | tataaaacgg | aagcagcctc | 600  |
| togatataac ctgacgatct cagacgtcag                         | cgtgagtgat | gtgccatttc | ctttctctgc | 660  |
| ccagtctggg gctggggtgc caggctgggg                         | catcgcgctg | ctggtgctgg | tctgtgttct | 720  |
| ggttgcgctg gccattgtct atctcattgc                         | cttggctgtc | tgtcagtgcc | gccgaaagaa | 780  |
| ctacgggcag ctggacatct ttccagcccg                         | ggatacctac | catcctatga | gcgagtaccc | 840  |
| cacctaccac acccatgggc gctatgtgcc                         | ccctagcagt | accgatcgta | gcccctatga | 900  |
| gaaggtttct gcaggtaatg gtggcagcag                         | cctctcttac | acaaacccag | cagtggcagc | 960  |
| cacttctgcc aacttgtagg ggcacgtcgc                         | ccgctgagct | gagtggccag | ccagtgccat | 1020 |
| tccactccac tcaggttctt cagggccaga                         | gcccctgcac | cctgtttggg | ctggtgagct | 1080 |
| gggagttcag gtgggctgct cacagcctcc                         | ttcagaggcc | ccaccaattt | ctcggacact | 1140 |
| tctcagtgtg tggaagctca tgtgggcccc                         | tgagggctca | tgcctgggaa | gtgttgtggt | 1200 |
| gggggctccc aggaggactg gcccagagag                         | ccctgagata | gcggggatcc | tgaactggac | 1260 |
| tgaataaaac gtggtctccc actgc                              |            |            |            | 1285 |

<210> 67 <211> 1517 <212> DNA <213> Homo sapien

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| <211> 524<br><212> DNA<br><213> Homo | o sapien     |              |            |            |            |     |
|--------------------------------------|--------------|--------------|------------|------------|------------|-----|
| <400> 68<br>gccctgatca               | gageceeeeg   | gtagaaggca   | ctccatggcc | tgccataacc | tcctatctcc | 60  |
| ccaggctgtc                           | tgtcagtgcc   | gccgaaagaa   | ctacgggcag | ctggacatct | ttccagcccg | 120 |
| ggatacctac                           | catcctatga   | gcgagtaccc   | cacctaccac | acccatgggc | gctatgtgcc | 180 |
| ccctagcagt                           | accgatcgta   | gcccctatga   | gaaggtgaga | ttgggcccca | caggccaggg | 240 |
| gaagcagagg                           | gtttggctgg   | gcaaggattc   | tgaagggggt | acttggaaaa | cccaaagagc | 300 |
| ttggaagagg                           | tgagaagtgg   | cgtgaagtga   | gcaggggagg | gcctggcaag | gatgaggggc | 360 |
| agaggtcaga                           | ggagttttgg   | gggacaggcc   | tgggaggaga | ctatggaaga | aaggggccct | 420 |
| caagagggag                           | tggccccact   | gccagaattc   | ctaaaagatc | attggccgtc | cacattcatg | 480 |
| ctggctggcg                           | ctggctgaac   | tggtgccacc   | gtggcagttt | tgtt       |            | 524 |
|                                      |              |              |            |            |            |     |
|                                      | gagtagggag   | agggaaggct   | taagagggga | agaaatgcag | gggccatgag | 60  |
| ccaaggccta                           | tgggcagaga   | gaaggaggct   | gctgcaggga | aggaggcggc | caacccaggg | 120 |
| gttactgagg                           | ctgcccactc   | cccagtcctc   | ctggtattat | ttctctggtg | gccagagctt | 180 |
| atattttctt                           | cttgctctta   | tttttccttc   | ataaagaccc | aaccctatga | ctttaacttc | 240 |
| ttacagctac                           | cacageceet   | aaacccgcaa   | cagttgttac | aggttctggt | catgcaagct | 300 |
| ctaccccagg                           | g tggagaaaag | gagacttcgg   | ctacccagag | aagttcagtg | cccagctcta | 360 |
| ctgagaagaa                           | tgctgtgagt   | atgaccagca   | gcgtactctc | cagccacagc | cccggttcag | 420 |
| gctcctccac                           | c cactcaggga | caggatgtca   | ctctggcccc | ggccacggaa | ccagcttcag | 480 |
| gttcagctgo                           | c cacctgggga | caggatgtca   | cctcggtccc | agtcaccagg | ccagccctgg | 540 |
| gctccacca                            | c cccgccagcc | : cacgatgtca | cctcagcccc | ggacaacaag | ccagccccgg | 600 |
| gctccaccg                            | cccccagcc    | : cacggtgtca | cctcggcccc | ggacaccagg | ccggccccgg | 660 |
| gctccaccg                            | c ccccccagco | : catggtgtca | ceteggeeee | ggacaacagg | cccgccttgg | 720 |
| gctccaccg                            | c ccctccagtc | cacaatgtca   | cctcggcctc | aggctctgca | tcaggctcag | 780 |
| cttctactc                            | t ggtgcacaac | ggcacctctg   | ccagggctac | cacaacccca | gccagcaaga | 840 |
| gcactccat                            | t ctcaattccc | agccaccact   | ctgatactcc | taccaccctt | gccagccata | 900 |
| gcaccaaga                            | c tgatgccagt | : agcactcacc | atagcacggt | acctcctctc | acctcctcca | 960 |

| atcacagcac ttctccccag                                    | rtatataata | gggtgtgttt   | ctttttccta   | tcttttcaca | 1020 |
|--|------------|--------------|--------------|------------|------|
|  |            |              |              |            | 1080 |
| tttcaaacct ccagtttaat                                    |            |              |              |            | 1000 |
| tgcagagaga catttctgaa                                    | atgtttttgc | agatttataa   | acaagggggt   | tttctgggcc | 1140 |
| tctccaatat taagttcagg                                    | ccaggatctg | tggtggtaca   | attgactctg   | gccttccgag | 1200 |
| aaggtaccat caatgtccac                                    | gacgtggaga | cacagttcaa   | tcagtataaa   | acggaagcag | 1260 |
| cctctcgata taacctgacg                                    | atctcagacg | tcagcgtgag   | tgatgtgcca   | tttcctttct | 1320 |
| ctgcccagtc tggggctggg                                    | gtgccaggct | ggggcatcgc   | gctgctggtg   | ctggtctgtg | 1380 |
| ttctggttgc gctggccatt                                    | gtctatctca | ttgccttggc   | tgtctgtcag   | tgccgccgaa | 1440 |
| agaactacgg gcagctggac                                    | atctttccag | cccgggatac   | ctaccatcct   | atgagcgagt | 1500 |
| accccaccta ccacacccat                                    | gggcgctatg | tgccccctag   | cagtaccgat   | cgtagcccct | 1560 |
| atgagaaggt ttctgcaggt                                    | aatggtggca | gcagcctctc   | ttacacaaac   | ccagcagtgg | 1620 |
| cagecaette tgecaacttg                                    | taggggcacg | togcoogatg   | agctgagtgg   | ccagccagtg | 1680 |
| ccattccact ccactcaggt                                    | tcttcagggc | cagagcccct   | gcaccctgtt   | tgggctggtg | 1740 |
| agctgggagt tcaggtgggc                                    | tgctcacagc | ctccttcaga   | ggccccacca   | atttctcgga | 1800 |
| cacttctcag tgtgtggaag                                    | ctcatgtggg | cccctgaggg   | ctcatgcctg   | ggaagtgttg | 1860 |
| tggtggggc tcccaggagg                                     | actggcccag | agagccctga   | gatagcgggg   | atcctgaact | 1920 |
| ggactgaata aaacgtggtc                                    | tcccactgc  |              |              |            | 1949 |
| <210> 70<br><211> 1803<br><212> DNA<br><213> Homo sapien |            |              |              |            |      |
| <400> 70<br>ggtagcgcaa gcagaacaca                        | gaccagcacc | agcagcgcga   | tgccccagcc   | gggcacccag | 60   |
| teteetttet teetgetget                                    | gctcctcaca | gtgcttacag   | ctaccacago   | ccctaaaccc | 120  |
| gcaacagttg ttacaggttc                                    | tggtcatgca | agetetacee   | caggtggaga   | aaaggagact | 180  |
| tcggctaccc agagaagttc                                    | agtgcccagc | tctactgaga   | . agaatgctgt | gagtatgacc | 240  |
| agcagcgtac tetecageca                                    | cageceeggt | tcaggctcct   | ccaccactca   | gggacaggat | 300  |
| gteactetgg ceceggecae                                    | ggaaccagct | tcaggttcag   | ctgccacctg   | gggacaggat | 360  |
| gtcacctcgg tcccagtcac                                    | caggccagcc | : ctgggctcca | ccaccccgc    | agcccacgat | 420  |

gtcacctcag ccccggacaa caagccagcc ccgggctcca ccgcccccc agcccacggt

gtcacctcgg ccccggacac caggccggcc ccgggctcca ccgcccccc agcccatggt

gtcacctcgg ccccggacaa caggcccgcc ttgggctcca ccgcccctcc agtccacaat

480

540

| gtcacctcgg cctcaggctc tgcatcaggc to                           | cagcttcta ctctggtgca caacggcacc 6  | 60  |
|---|------------------------------------|-----|
| tctgccaggg ctaccacaac cccagccagc as                           | agagcactc cattetcaat teecagecac 7  | 20  |
| cactetgata etectaceae cettgecage ea                           | atagcacca agactgatgc cagtagcact 7  | 80  |
| caccatagca eggtacetee teteacetee te                           | ccaatcaca gcacttctcc ccagttgtct 8  | 40  |
| actggggtct ctttcttttt cctgtctttt ca                           | acatttcaa acctccagtt taattcctct 9  | 00  |
| ctggaagatc ccagcaccga ctactaccaa ga                           | agctgcaga gagacatttc tgaaatgttt 9  | 60  |
| ttgcagattt ataaacaagg gggttttctg gg                           | geeteteea atattaagtt caggeeagga 10 | 20  |
| tctgtggtgg tacaattgac tctggccttc cg                           | gagaaggta ccatcaatgt ccacgacgtg 10 | 80  |
| gagacacagt tcaatcagta taaaacggaa go                           | cageetete gatataaeet gaegatetea 11 | 40  |
| gacgtcagcg tgagtgatgt gccatttcct tt                           | tetetgeee agtetgggge tggggtgeea 12 | 00  |
| ggctggggca tcgcgctgct ggtgctggtc tg                           | gtgttctgg ttgcgctggc cattgtctat 12 | 60  |
| ctcattgcct tggctgtctg tcagtgccgc cg                           | gaaagaact acgggcagct ggacatcttt 13 | 20  |
| ccagcccggg atacctacca tcctatgagc ga                           | agtacecca ectaceacae ecatgggege 13 | 80  |
| tatgtgcccc ctagcagtac cgatcgtagc co                           | cctatgaga aggtttctgc aggtaatggt 14 | 40  |
| ggcagcagcc tetettacac aaacccagca g                            | tggcagcca cttctgccaa cttgtagggg 15 | 00  |
| cacgtcgccc gctgagctga gtggccagcc ag                           | gtgccattc cactccactc aggttcttca 15 | 60  |
| gggccagagc ccctgcaccc tgtttgggct gg                           | gtgagctgg gagttcaggt gggctgctca 16 | 20  |
| cagecteett cagaggeece accaatttet eg                           | ggacacttc tcagtgtgtg gaagctcatg 16 | 80  |
| tgggcccctg agggctcatg cctgggaagt g                            | ttgtggtgg gggctcccag gaggactggc 17 | 740 |
| ccagagagcc ctgagatagc ggggatcctg as                           | actggactg aataaaacgt ggtctcccac 18 | 300 |
| tgc   | 18                                 | 303 |
| <210> 71<br><211> 1258<br><212> DNA<br><213> Homo sapien      |                                    |     |
| <pre>&lt;400&gt; 71 taggaggtag gggagggggc ggggttttgt ca</pre> | acctgtcac ctgctccggc tgtgctatgg    | 60  |
| gcgggcgggc ggggagtggg gggaccggta ta                           | aaageggta ggegeetgtg ceegeteeae 1  | L20 |
| ctctcaagca gccagcgcct gcctgaatct g                            | ttctgcccc ctccccaccc atttcaccac 1  | 180 |
| caccatgaca ccgggcaccc agtctccttt c                            | ttcctgctg ctgctcctca cagtgcttac 2  | 240 |
|   |                                    |     |

agttgttaca ggttctggtc atgcaagctc taccccaggt ggagaaaagg agacttcggc

tacccagaga agttcagtgc ccagctctac tgagaagaat gcttttaatt cctctctgga

300

PCT/US03/18934 WO 03/106648

| agateccage | accgactact | accaagagct | gcagagagac | atttctgaaa | tgtttttgca | 420  |
|------------|------------|------------|------------|------------|------------|------|
| gatttataaa | caagggggtt | ttctgggcct | ctccaatatt | aagttcaggc | caggatctgt | 480  |
| ggtggtacaa | ttgactctgg | ccttccgaga | aggtaccatc | aatgtccacg | acgtggagac | 540  |
| acagttcaat | cagtataaaa | cggaagcagc | ctctcgatat | aacctgacga | tctcagacgt | 600  |
| cagcgtgagt | gatgtgccat | ttcctttctc | tgcccagtct | ggggctgggg | tgccaggctg | 660  |
| gggcatcgcg | ctgctggtgc | tggtctgtgt | tctggttgcg | ctggccattg | tctatctcat | 720  |
| tgccttggct | gtctgtcagt | gccgccgaaa | gaactacggg | cagctggaca | tctttccagc | 780  |
| ccgggatacc | taccatccta | tgagcgagta | cccacctac  | cacacccatg | ggcgctatgt | 840  |
| gccccctagc | agtaccgatc | gtageceeta | tgagaaggtt | tctgcaggta | atggtggcag | 900  |
| cagcetetet | tacacaaacc | cagcagtggc | agccacttct | gccaacttgt | aggggcacgt | 960  |
| cgcccgctga | gctgagtggc | cagccagtgc | cattccactc | cactcaggtt | cttcagggcc | 1020 |
| agagcccctg | caccctgttt | gggctggtga | gctgggagtt | caggtgggct | gctcacagcc | 1080 |
| tccttcagag | gccccaccaa | tttctcggac | acttctcagt | gtgtggaagc | tcatgtgggc | 1140 |
| ccctgagggc | tcatgcctgg | gaagtgttgt | ggtgggggct | cccaggagga | ctggcccaga | 1200 |
| gagccctgag | atagcgggga | tcctgaactg | gactgaataa | aacgtggtct | cccactgc   | 1258 |

<210> 72

<211> 2045 <212> DNA <213> Homo sapien

<400> 72

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|                         |             |              | _            |            |              | 780  |
|-------------------------|-------------|--------------|--------------|------------|--------------|------|
|                         | ccagtccaca  |              |              |            |              |      |
| tactctggtg              | cacaacggca  | cctctgccag   | ggctaccaca   | accccagcca | gcaagagcac   | 840  |
| tccattctca              | attcccagcc  | accactctga   | tactcctacc   | acccttgcca | gccatagcac   | 900  |
| caagactgat              | gccagtagca  | ctcaccatag   | cacggtacct   | cctctcacct | cctccaatca   | 960  |
| cagcacttct              | ccccagttgt  | ctactggggt   | ctctttcttt   | ttcctgtctt | ttcacatttc   | 1020 |
| aaacctccag              | tttaattcct  | ctctggaaga   | tcccagcacc   | gactactacc | aagagctgca   | 1080 |
| gagagacatt              | tctgaaatgg  | tgagtatcgg   | cctttccttc   | cccatgctcc | cctgaagcag   | 1140 |
| ccatcagaac              | tgtccacacc  | ctttgcatca   | agcctgagtc   | ctttccctct | caccccagtt   | 1200 |
| ttttgcagat              | ttataaacaa  | gggggtttc    | tgggcctctc   | caatattaag | ttcaggccag   | 1260 |
| gatctgtggt              | ggtacaattg  | actctggcct   | tccgagaagg   | taccatcaat | gtccacgacg   | 1320 |
| tggagacaca              | gttcaatcag  | tataaaacgg   | aagcagcctc   | tcgatataac | ctgacgatct   | 1380 |
| cagacgtcag              | cgtgagtgat  | gtgccatttc   | ctttctctgc   | ccagtctggg | gctggggtgc   | 1440 |
| caggctgggg              | categegetg  | ctggtgctgg   | tctgtgttct   | ggttgcgctg | gccattgtct   | 1500 |
| atctcattgc              | cttggctgtc  | tgtcagtgcc   | gccgaaagaa   | ctacgggcag | ctggacatct   | 1560 |
| ttccagcccg              | ggatacctac  | catcctatga   | gcgagtaccc   | cacctaccac | acccatgggc   | 1620 |
| gctatgtgcc              | ccctagcagt  | accgatcgta   | gcccctatga   | gaaggtttct | gcaggtaatg   | 1680 |
| gtggcagcag              | cctctcttac  | acaaacccag   | cagtggcagc   | cacttctgcc | aacttgtagg   | 1740 |
| ggcacgtcgc              | ccgctgagct  | gagtggccag   | ccagtgccat   | tccactccac | tcaggttctt   | 1800 |
| cagggccaga              | gcccctgcac  | cctgtttggg   | ctggtgagct   | gggagttcag | gtgggctgct   | 1860 |
| cacageetee              | ttcagaggcc  | ccaccaattt   | ctcggacact   | tctcagtgtg | tggaagctca   | 1920 |
| tgtgggccc               | tgagggctca  | tgcctgggaa   | gtgttgtggt   | gggggctccc | aggaggactg   | 1980 |
| gcccagagag              | ccctgagata  | geggggatee   | tgaactggac   | tgaataaaac | gtggteteec   | 2040 |
| actgc                   |             |              |              |            |              | 2045 |
| <210> 73 <211> 126      |             |              |              |            |              |      |
| <212> DNA<br><213> Horr | o sapien    |              |              |            |              |      |
| <400> 73<br>taggaggtag  | 1 333ø33330 | : ggggttttgt | : cacctgtcac | ctgctccggc | : tgtgctatgg | 60   |

gcgggcggc gggagtggg gggaccggta taaagcggta ggcgcctgtg cccgctccac

ctctcaagca gccagcgcct gcctgaatct gttctgcccc ctccccaccc atttcaccac

caccatgaca cogggcacco agtotocttt cttcctgctg ctgctcctca cagtgcttac

120 180

| agctaccaca | gcccctaaac | ccgcaacagt | tgttacaggt | tctggtcatg | caagctctac | 300  |
|------------|------------|------------|------------|------------|------------|------|
| cccaggtgga | gaaaaggaga | cttcggctac | ccagagaagt | tcagtgccca | gctctactga | 360  |
| gaagaatgct | atcccagcac | cgactactac | caagagctgc | agagagacat | ttctgaaatg | 420  |
| tttttgcaga | tttataaaca | agggggtttt | ctgggcctct | ccaatattaa | gttcaggcca | 480  |
| ggatctgtgg | tggtacaatt | gactctggcc | ttccgagaag | gtaccatcaa | tgtccacgac | 540  |
| gtggagacac | agttcaatca | gtataaaacg | gaagcagcct | ctcgatataa | cctgacgatc | 600  |
| tcagacgtca | gcgtgagtga | tgtgccattt | cctttctctg | cccagtctgg | ggctggggtg | 660  |
| ccaggctggg | gcatcgcgct | gctggtgctg | gtctgtgttc | tggttgcgct | ggccattgtc | 720  |
| tatctcattg | ccttggctgt | ctgtcagtgc | cgccgaaaga | actacgggca | gctggacatc | 780  |
| tttccagccc | gggataccta | ccatcctatg | agcgagtacc | ccacctacca | cacccatggg | 840  |
| cgctatgtgc | cccctagcag | taccgatcgt | agcccctatg | agaaggtttc | tgcaggtaat | 900  |
| ggtggcagca | gcctctctta | cacaaaccca | gcagtggcag | ccacttctgc | caacttgtag | 960  |
| gggcacgtcg | cccgctgagc | tgagtggcca | gccagtgcca | ttccactcca | ctcaggttct | 1020 |
| tcagggccag | agcccctgca | ccctgtttgg | gctggtgagc | tgggagttca | ggtgggctgc | 1080 |
| tcacagcctc | cttcagaggc | cccaccaatt | tctcggacac | ttctcagtgt | gtggaagctc | 1140 |
| atgtgggccc | ctgagggctc | atgcctggga | agtgttgtgg | tgggggctcc | caggaggact | 1200 |
| ggcccagaga | gccctgagat | agcggggatc | ctgaactgga | ctgaataaaa | cgtggtctcc | 1260 |
| cactgc     |            |            |            |            |            | 1266 |

<210> 74

<211> 1189

<212> DNA

<213> Homo sapien

<400> 74 taggaggtag gggaggggc ggggttttgt cacctgtcac ctgctccggc tgtgctatgg 60 gcgggcgggc ggggagtggg gggaccggta taaagcggta ggcgcctgtg cccgctccac 120 180 ctctcaagca gccagcgcct gcctgaatct gttctgcccc ctccccaccc atttcaccac caccatgaca ccgggcaccc agtctccttt cttcctgctg ctgctcctca cagtgcttac 240 300 agttgttaca ggttctggtc atgcaagctc taccccaggt ggagaaaagg agacttcggc tacccagaga agttcagtgc ccagctctac tgagaagaat gcttttttgc agatttataa 360 acaagggggt tttctgggcc tctccaatat taagttcagg ccaggatctg tggtggtaca 420 480 attgactctg gccttccgag aaggtaccat caatgtccac gacgtggaga cacagttcaa tcagtataaa acggaagcag cctctcgata taacctgacg atctcagacg tcagcgtgag 540

| tgatgtgcca | tttcctttct | ctgcccagtc | tggggctggg | gtgccaggct | ggggcatcgc | 600  |
|------------|------------|------------|------------|------------|------------|------|
| gctgctggtg | ctggtctgtg | ttctggttgc | gctggccatt | gtctatctca | ttgccttggc | 660  |
| tgtctgtcag | tgccgccgaa | agaactacgg | gcagctggac | atctttccag | cccgggatac | 720  |
| ctaccatcct | atgagcgagt | accccaccta | ccacacccat | gggcgctatg | tgccccctag | 780  |
| cagtaccgat | cgtagcccct | atgagaaggt | ttctgcaggt | aatggtggca | gcagcctctc | 840  |
| ttacacaaac | ccagcagtgg | cagccacttc | tgccaacttg | taggggcacg | tcgcccgctg | 900  |
| agctgagtgg | ccagccagtg | ccattccact | ccactcaggt | tcttcagggc | cagageeeet | 960  |
| gcaccctgtt | tgggctggtg | agctgggagt | tcaggtgggc | tgctcacagc | ctccttcaga | 1020 |
| ggccccacca | atttctcgga | cacttctcag | tgtgtggaag | ctcatgtggg | cccctgaggg | 1080 |
| ctcatgcctg | ggaagtgttg | tggtggggc  | tcccaggagg | actggcccag | agagccctga | 1140 |
| gatagcgggg | atcctgaact | ggactgaata | aaacgtggtc | tcccactgc  |            | 1189 |
|            |            |            |            |            |            |      |

<210> 75

<211> 1216

<212> DNA

<213> Homo sapien

<400> 75

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| ctcaggttct  | tcagggccag | agcccctgca   | ccctgtttgg | gctggtgagc | tgggagttca   | 1020 |
|---|------------|--------------|------------|------------|--------------|------|
| ggtgggctgc  | tcacageete | cttcagaggc   | cccaccaatt | tctcggacac | ttctcagtgt   | 1080 |
| gtggaagctc  | atgtgggccc | ctgagggctc   | atgcctggga | agtgttgtgg | tgggggctcc   | 1140 |
| caggaggact  | ggcccagaga | gccctgagat   | agcggggatc | ctgaactgga | ctgaataaaa   | 1200 |
| cgtggtctcc  | cactgc     |              |            |            |              | 1216 |
| <210> 76<br><211> 2090<br><212> DNA<br><213> Homo | sapien     |              |            |            |              |      |
| <400> 76  | aaaaaaaaac | ggggttttgt   | cacctqtcac | ctactccggc | tgtgctatgg   | 60   |
|   |            | gggaccggta   |            |            |              | 120  |
|   |            |              |            |            |              | 180  |
|   |            | gcctgaatct   |            |            |              | 240  |
|   |            | agtctccttt   |            |            |              |      |
|   |            | ccgcaacagt   |            |            |              | 300  |
| cccaggtgga  | gaaaaggaga | cttcggctac   | ccagagaagt | tcagtgccca | gctctactga   | 360  |
| gaagaatgct  | gtgagtatga | ccagcagcgt   | actctccagc | cacagccccg | gttcaggctc   | 420  |
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| tccattctca | attcccagcc   | accactctga | tactcctacc | acccttgcca | gccatagcac | 900    |
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<212> DNA

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| caccgcccc ccagcccacg gtgtcacctc ggccccggac accaggccgg ccccgggc            | tc 660   |
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| caccgcccct ccagtccaca atgtcacctc ggcctcaggc tctgcatcag gctcagct           | tc 780   |
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| gagagacatt tetgaaatgt ttttgeagat ttataaacaa gggggtttte tgggeete           | ctc 1140 |
| caatattaag ttcaggccag gatctgtggt ggtacaattg actctggcct tccgaga            | agg 1200 |
| taccatcaat gtccacgacg tggagacaca gttcaatcag tataaaacgg aagcagc            | ctc 1260 |
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| cttctgccaa cttgtagggg cacgtcgccc gctgagctga                               | ttc 1560 |
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| caccatgaca ccgggcaccc agtctccttt cttcctgctg ctgctcctca cagtgct            | ttac 240 |
|   |          |

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| gaagaatgct                          | gtgagtatga       | ccagcagcgt   | actctccagc   | cacageeeeg  | gttcaggctc        | 420  |
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| caagactgat                          | gccagtagca       | ctcaccatag   | cacggtacct   | cctctcacct  | cctccaatca        | 960  |
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| catctttcca                          | gcccgggata       | cctaccatco   | tatgagcgag   | taccccacct  | accacaccca        | 1200 |
| tgggcgctat                          | gtgcccccta       | gcagtaccga   | tegtageece   | tatgagaagg  | tttctgcagg        | 1260 |
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|                                     |                  |              |              |             | a cctaccatcc      | 180  |
|                                     |                  |              |              |             | a gcagtaccga      | 240  |
|                                     |                  |              |              |             | t cttacacaaa      | 300  |

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| gggaagtgtt             | gtggtggggg | ctcccaggag | gactggccca | gagagccctg | agatagcggg | 600 |
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| cccctatgag             | aaggtttctg | caggtaatgg | tggcagcagc | ctctcttaca | caaacccagc | 540 |
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| cagtgccatt             | ccactccact | caggttcttc | agggccagag | cccctgcacc | ctgtttgggc | 660 |
| tggtgagctg             | ggagttcagg | tgggctgctc | acagcctcct | tcagaggccc | caccaatttc | 720 |
| tcggacactt             | ctcagtgtgt | ggaagctcat | gtgggcccct | gagggctcat | gcctgggaag | 780 |
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| gaactggact             | gaataaaacg | tggtctccca | ctgc       |            |            | 874 |
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| <400> 82<br>ttttgctttt | ttgcacccag | aggcaaaatg | ggtggagcac | tatgcccagg | ggagcccttc | 60  |
| ccgaggagtc             | ccaggggtga | gcctctgtgc | ccctaatcat | ctcctaggaa | tggagggtag | 120 |

79

| accgagaaag gct                                       | ggcatag gg  | ggaggttt  | cccaggtaga | agaagaagtg | tcagcagacc | 180  |
|--|-------------|-----------|------------|------------|------------|------|
| aggtgagcgt ggg                                       | tgccagt gg  | ggttcttg  | ggagcttcaa | ggaagcaagg | aacgctccct | 240  |
| cetteetete etg                                       | gtctttc tc  | tatgggac  | ctagtaaata | attactgcag | ccacctgagg | 300  |
| ctggaaaacc act                                       | ccaggtg gg  | ggaggaga  | gagtttagtt | ttcttgctcc | tattttcctc | 360  |
| ctcctggaga cct                                       | ccctctc tc  | ggctttac  | aaagacacag | atacaccccg | cccccaaac  | 420  |
| acacacacac aca                                       | cacacac ac  | acctcctt  | aggctggaac | agcagagaat | ggagggacaa | 480  |
| gggggctgat tag                                       | agccaag aa  | gagggagt  | gaaggagagc | agagggagga | gggcagccct | 540  |
| gtttacagtc acc                                       | tggctgg tg  | gggtggca  | ggtgctctct | ctgaattaac | cctttgagag | 600  |
| ctggccagga ctc                                       | tggactg at  | taccccag  | cctggggtgg | catccagggg | ctctaggagg | 660  |
| taccttttgc tcc                                       | tcaccct gg  | gatctcttt | tccttccacc | caggtttctg | caggtaatgg | 720  |
| tggcagcagc ctc                                       | tcttaca ca  | aacccagc  | agtggcagcc | acttctgcca | acttgtaggg | 780  |
| gcacgtcgcc cgc                                       | tgagctg ag  | gtggccagc | cagtgccatt | ccactccact | caggttcttc | 840  |
| agggccagag ccc                                       | ctgcacc ct  | gtttgggc  | tggtgagctg | ggagttcagg | tgggctgctc | 900  |
| acagectect tea                                       | gaggccc ca  | accaatttc | tcggacactt | ctcagtgtgt | ggaagctcat | 960  |
| gtgggcccct gag                                       | ggctcat go  | cctgggaag | tgttgtggtg | ggggctccca | ggaggactgg | 1020 |
| cccagagagc cct                                       | gagatag cg  | gggateet  | gaactggact | gaataaaacg | tggtctccca | 1080 |
| ctgc   |             |           |            |            |            | 1084 |
| <210> 83<br><211> 1194<br><212> DNA<br><213> Homo sa | pien        |           |            |            |            |      |
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| gegggeggge ggg                                       | gagtggg gg  | ggaccggta | taaagcggta | ggcgcctgtg | cccgctccac | 120  |
| ctctcaagca gcc                                       | cagegeet go | cctgaatct | gttctgcccc | ctccccaccc | atttcaccac | 180  |
| caccatgaca ccg                                       | ggcaccc ag  | gtctccttt | cttcctgctg | ctgctcctca | cagtgcttac | 240  |
| agctaccaca gco                                       | cctaaac c   | cgcaacagt | tgttacaggt | tctggtcatg | caagctctac | 300  |
| cccaggtgga gaa                                       | aaaggaga c  | ttcggctac | ccagagaagt | tcagtgccca | gctctactga | 360  |
| gaagaatgct gtg                                       | gagtatga c  | cagcagcgt | actctccagc | cacageeeeg | gttcaggctc | 420  |
| ctccaccact cag                                       | ggacagg a   | tgtcactct | ggccccggcc | acggaaccag | cttcaggttc | 480  |
| agctgccacc tgg                                       | gggacagg a  | tgtcacctc | ggtcccagtc | accaggccag | ccctgggctc | 540  |

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80

| caccgcccc ccagcccatg gtgtcacctc ggccccggac aacaggcccg ccttgggctc 7   | 720 |
|--|-----|
|  |     |
| caccgccct ccagtccaca atgtcacctc ggcctcaggc tctgcatcag gctcagcttc 7   | 780 |
| tactetggtg cacaacggca cetetgecag ggctaccaca accecageca gcaagagcac    | 840 |
| tccattctca attcccagcc accactctga tactcctacc acccttgcca gccatagcac    | 900 |
| caagactgat gccagtagca ctcaccatag cacggtacct cctctcacct cctccaatca    | 960 |
| cagcacttct ccccagttgt ctactggggt ctctttcttt ttcctgtctt ttcacatttc 10 | 020 |
| aaacetecag tttaatteet etetggaaga teecageace gaetaetaee aagagetgea 10 | 080 |
| gagagacatt totgaaatgt ttttgcagat ttataaacaa gggggttttc tgggcototo 13 | 140 |
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<211> 2623

<212> DNA

<213> Homo sapien

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| agggcacctc tas | aggateeg a | aggcacctgc | agtacgtggc | agatgtgaac | gagtccaacg   | 1080 |
|----------------|------------|------------|------------|------------|--------------|------|
| tgtacgtggt gad | cgcagggc ( | cgcaagctct | acgggatgcc | cactgacttc | ggtttctgtg   | 1140 |
| tcaagcccaa caa | agcttcga a | aatggccaca | aggggcttcg | gatettetge | agtgaagatg   | 1200 |
| agcagagccg ca  | cctgctgg ( | etggctgcct | teegeetett | caagtacggg | gtgcagctgt   | 1260 |
| acaagaatta cc  | agcaggca ( | cagtctcgcc | atctgcatcc | atcttgtttg | ggctccccac   | 1320 |
| ccttgagaag tg  | cctcagat a | aataccctgg | tggccatgga | cttctctggc | catgctgggc   | 1380 |
| gtgtcattga ga  | accccgg (  | gaggctctga | gtgtggccct | ggaggaggcc | caggcctgga   | 1440 |
| ggaagaagac aa  | accaccgc   | ctcagcctgc | ccatgccagc | ctccggcacg | agcctcagtg   | 1500 |
| cagccatcca cc  | gcacccaa   | ctctggttcc | acgggcgcat | ttcccgtgag | gagagccagc   | 1560 |
| ggcttattgg ac  | agcagggc   | ttggtagacg | gcctgttcct | ggtccgggag | agtcagcgga   | 1620 |
| acccccaggg ct  | ttgtcctc   | tetttgtgcc | acctgcagaa | agtgaagcat | tatctcatcc   | 1680 |
| tgccgagcga gg  | aggagggc   | cgcctgtact | tcagcatgga | tgatggccag | acccgcttca   | 1740 |
| ctgacctgct gc  | agctcgtg   | gagttccacc | agctgaaccg | cggcatcctg | ccgtgcttgc   | 1800 |
| tgcgccattg ct  | gcacgcgg   | gtggccctct | gaccaggccg | tggactggct | catgcctcag   | 1860 |
| cccgccttca gg  | getgeeege  | cgcccctcca | cccatccagt | ggactctggg | gcgcggccac   | 1920 |
| aggggacggg at  | gaggagcg   | ggagggttcc | gccactccag | ttttctcctc | tgcttctttg   | 1980 |
| cctccctcag at  | agaaaaca   | gcccccactc | cagtccactc | ctgacccctc | tcctcaaggg   | 2040 |
| aaggccttgg gt  | ggccccct   | ctccttctcc | tagctctgga | ggtgctgctc | tagggcaggg   | 2100 |
| aattatggga ga  | aagtggggg  | cagcccaggc | ggtttcacgc | cccacacttt | gtacagaccg   | 2160 |
| agaggccagt to  | gatctgctc  | tgttttatac | tagtgacaat | aaagattatt | ttttgataaa   | 2220 |
| aaactcagaa ct  | tatctcgtc  | gcgagtttga | taaaaagtgt | aaaaaaactg | gggggaactt   | 2280 |
| catagggggt ca  | aaacatctc  | gctgccggcg | gataggactt | ggctaaactt | cttccgagcg   | 2340 |
| ggccccgtaa g   | ggtggtatg  | ctgataaaaa | tgggggggg  | ccccctctc  | agggggccct   | 2400 |
| ccagaacctt t   | tgggggtgg  | ggtacccttg | ggtggttaac | tagtgaacto | : tttcctcaaa | 2460 |
| aggttgccgc c   | ccctgtgta  | ttgtcgacaa | ttttcttggg | gggegggee  | : gttttcttt  | 2520 |
| caccacgett t   | tgttttccc  | gggtggggaa | cccacccctg | gtgtgtgtg  | ccccccgtt    | 2580 |
| tattttgggc g   | ccctttttg  | tggggggaaa | ttcccccgct | ttt        |              | 2623 |

<sup>&</sup>lt;210> 85 <211> 1036 <212> DNA <213> Homo sapien

<sup>&</sup>lt;400> 85

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| ccaattttcg                       | tctccctccc                           | ccagccaagg | tctcccaggg | gtgcagggag | agcggagctg | 180  |
| ctcagagctt                       | ggccaggttc                           | taagtgtgct | cctgaaagca | ggtcacccct | gagatcctca | 240  |
| gggtggggca                       | cagaggggca                           | ccctagcagg | taaagggagg | ccacgggatg | gcggtgggca | 300  |
| getggeette                       | tagtaacgag                           | ccctcagtgc | cttctgtgcc | tggggtccct | gccgacggga | 360  |
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| tgccttgtgt                       | gtggtgggcc                           | tggggctggc | gccgcagtct | ctgaacctgt | gtgacgcctg | 480  |
| cagggctggg                       | acctgacggt                           | gaagatgctg | gcgggcaacg | aattccaggt | gtccctgagc | 540  |
| agctccatgt                       | cggtgtcaga                           | gctgaaggcg | cagatcaccc | agaagatcgg | cgtgcacgcc | 600  |
| ttccagcagc                       | gtctggctgt                           | ccacccgagc | ggtgtggcgc | tgcaggacag | ggtccccctt | 660  |
| gccagccagg                       | gcctgggccc                           | cggcagcacg | gtcctgctgg | tggtggacaa | atgcgacgaa | 720  |
| cctctgagca                       | tcctggtgag                           | gaataacaag | ggccgcagca | gcacctacga | ggtgcggctg | 780  |
| acgcagaccg                       | tggcccacct                           | gaagcagcaa | gtgagcgggc | tggagggtgt | gcaggacgac | 840  |
| ctgttctggc                       | tgaccttcga                           | ggggaagccc | ctggaggacc | agctcccgct | gggggagtac | 900  |
| ggcctcaagc                       | ccctgagcac                           | cgtgttcatg | aatctgcgcc | tgcggggagg | cggcacagag | 960  |
| cctggcgggc                       | ggagctaagg                           | gccccaccag | catccgagca | ggatcaaggg | ccggaaataa | 1020 |
| aggctgttgt                       | aaagag                               |            |            |            |            | 1036 |
| <220><br><221> mise<br><222> (16 | c_feature<br>B)(208)<br>, c, g, or t | =          |            |            | •          |      |
| <400> 86<br>gctgcctcta           | taggtgctgg                           | tatataagta | ttatcgacat | catttaagta | atgatttaga | 60   |
|                                  | aaaaaaaatt                           |            |            |            |            | 120  |
|                                  | aaacttggaa                           | ·          |            |            |            | 180  |
|                                  | nnnnnnnnn                            |            |            |            |            | 240  |
|                                  | cttcaggacc                           |            |            |            | _          | 300  |
|                                  |                                      |            |            |            |            |      |

ttatctgctt tctgcaatca cgtctcttcc atggggcact gagcagagaa tggtgtggcc

PCT/US03/18934

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<223> n=a, c, g, or t

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<212> DNA

<213> Homo sapien

<400> 89

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| ctaccctact ccaccttca                                     | g cccctcccgc  | gggggtagcg   | cctctcattc   | ctgatgtctc   | 600  |
|--|---------------|--------------|--------------|--------------|------|
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| agatcatcag aggctgaaa                                     | t ctagaacttc  | atcccgggca   | atgaggttct   | cacagaaggt   | 720  |
| gcagttttat aactaacta                                     | c gtccacttat  | atatattcac   | actctacata   | tatatatata   | 780  |
| tatatatata tatatatat                                     | a tatatataca  | cacaaatgca   | cagtccccct   | cccactccgt   | 840  |
| tacctaactg tacgtcttt                                     | t catgtttata  | aactatacag   | aaaactgtat   | ttgctgaact   | 900  |
| aaggattgta ttggtgatt                                     | t ctagcaaaaa  | caaagtgata   | gaatttttgt   | ctagaatccc   | 960  |
| aaactggcaa cgatagtct                                     | c caagggacct  | ggccttgcca   | agggcctggg   | gcaaggtgtc   | 1020 |
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| ccctgtcacc cctgctgt                                      | cg tagctgttgg | cttcagggtg   | agaagtgaga   | agcagcttat   | 1200 |
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| ttgggatctg gaaaccag                                      | ca ggccaggcag | catccacagt   | gttagtccaa   | agggtcggac   | 1320 |
| cgtgtcgtca gcctagcg                                      | tt tggtcagtga | cggcctggac   | gggccaagga   | gactccgggc   | 1380 |
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| tggccctccc agccgtcc                                      | tg gggatgctca | gggtgcaggc   | agaggetegg   | gaggccggac   | 1620 |
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| caaagccggc cgtgctgc                                      | tg geeggggge  | caggtgggta   | caagctcctt   | tgtgctttgc   | 1740 |
| acaaacctga atccccca                                      | cc agagaggatg | g tgtgtgagga | gccagaaacg   | ctgaatccaa   | 1800 |
| ttaagagaga aaaataat                                      | aa taacaataa  | a tgatcttgga | a caagaaaaaa | aaaa         | 1854 |
| <210> 90<br><211> 1759<br><212> DNA<br><213> Homo sapien |               |              |              |              |      |
| <400> 90<br>atgtgaaaag aaaatagt                          | ta tetgtgett  | g gtgttgtgt  | g ctctcctaaa | a gttaaccaga | 60   |
| cgtgaagcca aaaacato                                      | aa ctgggactg  | a caacacaaga | a aagattctt  | aactgaggtg   | 12   |
| gttaaatggc cctgaaaa                                      |               |              |              |              | 18   |
| gaggagga gcgagcga  | age gecaggtee | c ggcagggact | t cacttggag  | c tggcgtactt | 24   |

ggtgaccgcc ttggtgccct cggacacggc gtgcttggcc agctcgccgg gcagcagcag

| gcgcacggcc             | gtctggatct  | ccgggacgtg   | atggtggagc   | gcttgttgta   | gtgcgccagg   | 360  |
|------------------------|-------------|--------------|--------------|--------------|--------------|------|
| cgggacgcct             | ctcccgcgat  | gcgctcgaag   | atgtcgttga   | ggaaggagtt   | catgatgccc   | 420  |
| atggccttgc             | accagatgcc  | ggtgtcgggg   | tggacccgct   | tcagcacctt   | gtacacgtag   | 480  |
| atggagtagc             | tctccttgcg  | gctgcgcttg   | cgcttcttgc   | cgtctttctt   | ctgggctttg   | 540  |
| gtgacggctt             | tcttggagcc  | cttcttggga   | gccggcgcga   | actttgcagg   | ctcaggcatg   | 600  |
| gccagaccca             | agaccgacac  | cgacccccga   | gaacgcaagc   | agagcggtag   | gctcggggtc   | 660  |
| taccggaaac             | gactgtgtac  | ttacagaggc   | tgtgcgcatg   | acgctgcgtt   | atggttcgcg   | 720  |
| agttttccgc             | ggcgcgcaat  | gcgagggaga   | cgagattatg   | taaatgagtg   | gattctggct   | 780  |
| gagctatcct             | attggctatc  | gggacaaaat   | ttgcttgagc   | caatcaaagt   | geteegtgga   | 840  |
| caatcgccgt             | tctgtctata  | aaaaggtgaa   | gcagcggcgt   | tttcggcgac   | tttcccgatc   | 900  |
| gccaggcagg             | agtttctctc  | ggtgactact   | atcgctgtca   | tgtctggtcg   | tggcaagcaa   | 960  |
| ggaggcaagg             | cccgcgccaa  | ggccaagtcg   | cgctcgtccc   | gegetggeet   | tcagttcccg   | 1020 |
| gtagggcgag             | tgcatcgctt  | gctgcgcaaa   | ggcaactacg   | cggagcgagt   | gggggccggc   | 1080 |
| gcgcccgtct             | acatggctgc  | ggtcctcgag   | tatctgaccg   | ccgagatcct   | ggagctggcg   | 1140 |
| ggcaacgcgg             | ctcgggacaa  | caagaagacg   | cgcatcatcc   | ctcgtcacct   | ccagctggcc   | 1200 |
| atccgcaacg             | acgaggaact  | gaacaagctg   | ctgggcaaag   | tcaccatcgc   | ccagggcggc   | 1260 |
| gtcttgccta             | acatccaggo  | : cgtactgctc | cctaagaaga   | cggagagtca   | ccacaaggca   | 1320 |
| aagggcaagt             | gaggetgaeg  | teeggeecaa   | gtgggcccag   | cccggcccgc   | gtctcgaagg   | 1380 |
| ggcacctgtg             | aactcaaaag  | gctcttttca   | gagccaccca   | cgttttcaaa   | taaaagagtt   | 1440 |
| gttaatgctg             | gecaetetea  | gtccagcgtt   | cctcagtagt   | gaatagcgaa   | cctggagctg   | 1500 |
| acgggacggg             | acgggacggg  | g acgggacggg | geggggeggg   | gcggggcggg   | gtgtgtgtgt   | 1560 |
| gtgcgcgccg             | tettecatet  | ggagcacgta   | actgccttgg   | ctcttcgatg   | agtgggtccc   | 1620 |
| cagtcctagg             | acttcccago  | g gcaggtgcag | g gcaccaaacg | teetgggege   | : cgccacggtc | 1680 |
| cgctccacac             | : agtcacaaa | accagegee    | g cgggcagtac | : ccaacgcgct | gaagtgttgc   | 1740 |
| gcgcggagcg             | gegettee    |              |              |              |              | 1759 |
| .010                   | •           |              |              |              |              |      |
| <210> 91<br><211> 123  | 14          |              |              |              |              |      |
| <212> DNA              |             |              |              |              |              |      |
| <213> Hor              | no sapien   |              |              |              |              |      |
| <400> 91<br>ggtcactctc | c tactcaagt | t ctacttata  | t aacagcaats | g cagetetett | cataaagctg   | 60   |

gctgttgtgt agtttatgtt ggggaatcag ttcatggttt aaaaagttct gtcaatgcag

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| agaacaagcc             | ggtgtgtttt   | atggagaggc   | tgtttaatct | ccactgtgag   | acagtaaata | 180  |
|------------------------|--------------|--------------|------------|--------------|------------|------|
| tttggctgtt             | gcatcatcgt   | gaagcttatg   | atcacagtct | ggcgccatct   | ccatactaga | 240  |
| ctggagtctg             | atctgtcccg   | gcccagtgtc   | ctccaggaac | ctggcccctc   | atgcctccgt | 300  |
| gcttgcgcgt             | gtgccatttc   | ctctctccag   | aggacctttc | ctgcctagga   | ctcatcattg | 360  |
| tccccttcct             | ggtaagccat   | ccccgacctt   | ccaggcagaa | cctgctggct   | tctcctcagc | 420  |
| actttgcatg             | gatttcatgt   | cacagtcctg   | ggtgcactgt | gtcgccctct   | ctatgtgtca | 480  |
| gcctcccgtc             | ccctaccgtg   | ggctcctcca   | gggaggtgtg | gacattcatc   | ctcttccagg | 540  |
| cagccctcag             | gaatccaggg   | agaagataag   | gaggcggggc | gggcggaggg   | gggtgctcca | 600  |
| cacactcaga             | acactttcct   | ctgcacttac   | ttcattctgg | tttttcttt    | gggtccttgg | 660  |
| tgtttttaaa             | taaacccttt   | cctgtagttt   | geteceette | catggagggc   | tgtttcgagc | 720  |
| acagatetge             | tgggtgtctg   | tatttacaaa   | gagaaggggc | cactcgtgtg   | tgagcagcac | 780  |
| cgagggacag             | aggtaccttg   | cctgcttgtg   | tcccctccaa | gtccttctga   | tattttcctt | 840  |
| tccagctgtt             | gcctagtttc   | ctggtattaa   | ggagaatcaa | ctctctggat   | aaacgtggta | 900  |
| aatatggccc             | atagtcccat   | ctttttacag   | gcattttta  | cacctggagc   | agccagagga | 960  |
| cgcatgcatg             | gctcttcgga   | aggtaattta   | gggatcaccc | atgtaagttt   | cctaaggatt | 1020 |
| tctttaacat             | ggttcttctg   | attcagtccg   | gccaattaaa | tctaaatcca   | cccctgaaag | 1080 |
| ccatctggtg             | tggataacaa   | gcccacaaat   | gagcagtcag | ctttttgtgc   | cctttagggc | 1140 |
| ctgggacaac             | cacgggatct   | aaaaggggct   | ggaactagag | gtcttgagct   | cctgttccta | 1200 |
| aaatcatctt             | catcctatat   | ctgcagtctt   | ctcc       |              |            | 1234 |
|                        | •            |              |            |              |            |      |
| <210> 92<br><211> 730  | )            |              |            |              |            |      |
| <212> DNA<br><213> Hor | no sapien    |              |            |              |            |      |
| <400> 92               |              |              |            |              |            |      |
|                        | gagaaagaac   | tgactgaaac   | gtttgagata | tataggaaac   | atcaaaaggt | 60   |
| gataaaattt             | ccctagaato   | tccactatct   | caaagatgaa | gaaagttctc   | ctcctgatca | 120  |
| cagccatctt             | gggcagtggc   | : tgttggtttc | ccagtctctc | aagaccagga   | acgagaaaaa | 180  |
| agaagtgtaa             | gttacctttt   | ctcttttta    | catatcagtg | acagcgatga   | attagcttca | 240  |
| gggtttttt              | g tgttccctta | . cccatatcca | tttcgcccac | : ttccaccaat | tccatttcca | 300  |
| agatttccat             | ggtttagacg   | , taattttcct | attccaatac | ctgaatctgc   | ccctacaact | 360  |
| ccccttccta             | a gcgaaaagta | aacaagaagg   | aaaagtcacg | , ataaacctgg | tcacctgaaa | 420  |
| ttgaaattga             | a gccacttcct | : tgaagaatca | aaattcctgt | : taataaaaga | aaaacaaatg | 480  |

| · · · · · · · · · · · · · · · · · · ·                             | E40  |
|---|------|
| taattgaaat agcacacagc attototagt caatatottt agtgatotto tttaataaac | 540  |
| atgaaagcaa atcactaaag atattgacta gagaatgctg tgtgctattt caatatcttt | 600  |
| agtgatcttc tttaataaac atgaaagcat aaaaaaaaa agacgaaaaa aaaaggctgg  | 660  |
| gggcaccctg ggacaaagcg gtcccggggg ggattggttc ccggccaatt ccacaataag | 720  |
| ccgcacaaga  | 730  |
|   |      |
| <210> 93<br><211> 1159  |      |
| <212> DNA<br><213> Homo sapien                                    |      |
| -<br><400> 93   |      |
| ggggacagat ttctccattc cattatacct ttgagtatat aaaacagcta caatattcca | 60   |
| gggccagtca cttgccattt ctcataacag cgtcagagag aaagaactga ctgaaacgtt | 120  |
| tgagatatat aggaaacatc aaaaggtgat aaaatttccc tagaatctcc actatctcaa | 180  |
| agatgaagaa agttctcctc ctgatcacag ccatcttggc agtggctgtt ggtttcccag | 240  |
| tctctcaaga ccaggaacga gaaaaaagaa gtgtaagtta ccttttctct tttttacata | 300  |
| tcagtgacag cgatgaatta gcttcagggt tttttgtgtt cccttaccca tatccatttc | 360  |
| gcccacttcc accaattcca tttccaagat ttccatggtt tagacgtaat tttcctattc | 420  |
| caatacctga atctgcccct acaactcccc ttcctagcga aaagtaaaca agaaggaaaa | 480  |
| gtcacgataa acctggtcac ctgaaattga aattgagcca cttccttgaa gaatcaaaat | 540  |
| tcctgttaat aaaagaaaaa caaatgtaat tgaaatagca cacagcattc tctagtcaat | 600  |
| atctttagtg atcttcttta ataaacatga aagaagatca ctaaagatat tgactagaga | 660  |
| atgctgtgtg ctatttcaat tacatttgtt tttctttaat aaacatgaat tttgattctt | 720  |
| caaggaagtg gctcaatttc aatttcaggt gacctgaaat aaataacaga catatggtta | 780  |
| ttaattgcaa tgggtcattt tcttggaaac atatacattt tctgcatttt aatgacaact | 840  |
| attggcttaa aaatatatct agttcaagga ctgggaaacc atctgctcaa gatgtagaaa | 900  |
| gaaagcaaag gtctttagtg gtaagtagta gctgaaatat ttttttccta gaacagtcct | 960  |
| ctgggttcta atttaatctt agataagatt aaattatata tattaaatta taaattatta | 1020 |
| tagtagatta gatctatagt ctatagtata gattatattt cctcaattta tctagtaatt | 1080 |
| gacacaccat ccactttgtt tttgatgtga tgaaatgaca ggggccactg ttataggtga | 1140 |
| agcatgaagc ctttaaaat  | 1159 |
|   |      |

<sup>&</sup>lt;210> 94 <211> 1493

<212> DNA

<213> Homo sapien

<400> 94 ggagcccagc cgtgggattt tcaggtgttt tcatttggtg atcaggactg aacagagaga 60 actcaccatg gagtttgggc tgagctggct ttttcttgtg gctattttaa aaggtgtcca 120 gtgtgaggtg cagctgttgg agtctggggg aggcttggta cagcctgggg ggtccctgag 180 acteteetgt geageetetg gatteacett tageatetat gecatgaget gggteegeea 240 ggctccaggg aaggggctgg agtgggtcgc aagtatcagt ttcagtggtg gtagtacata 300 ctacgcagac tccgtgaagg gccgtttcac catctccaga gacaattcca agaccacgat 360 gcatctccac atgaacagcc tgagaaccga cgacacggcc gtctactact gtgcgaaacc 420 gtttccgtat tttgactact ggggccaggg aaccctggtc accgtctcga gtggcgatgg 480 gtccagtggc ggtagcgggg gcgcgtcgac tggcgaagtt gtgttgacgc agtttccagg 540 gcaccetgte tetgteteca ggggaaagag ccaccetete etgeagggee agteagagtg 600 cttagcagca gctacttagc ctggtatcag cagagacctg gccaggctcc caggctcctc 660 gtttatagtg catctgtgcg gcccaatgat attccagtca gggtccgtgg cagtgggtct 720 gggacagagt tcactctcac catcagcaga ctggtaacct gaagattttg cagtgtatta 780 ctgtcaacag ctatgggggc tcacctgacg tggactttcg ccccggggac caaggtggaa 840 gtccaaacga actgtggctg caccatctgt cttcatcttc ccgccatctg atgagcagtt 900 gaaatctgga actgeetetg ttgtgtgeet getgaataae ttetateeca gagaggeeaa 960 agtacagtgg aaggtggata acgccctacc aatcgggtaa ctcccaggag agtgtcacag 1020 agcaggacag cacaggacag acacctacag cctcagcagc accctgacgc tgagcaaagc 1080 agactacgag aaacacaaac tctacgcctg cgaagtcacc catcagggcc tgagctcgcc 1140 cgtcacaaag agcttcaaca ggggagagtg ttagagggag aagtgccccc acctgctcct 1200 cagttccage ctgaccccct cccatccttt ggcctctgac cctttttcca caggggacct 1260 acccctattg cggtcctcca gctcatcttt cacctcaccc ccctcctcct ccttggcttt 1320 aattatgcta atgttggagg gagcctgact aaataaagtg aatctttaaa acacaaaaaa 1380 aaggaaaaca aaaaaacaaa aaaaaaaaaa acacgcgggc ggacacccgg ggacaacggg 1440 gteccegggg teacactggt taccegteca attteccaca aaacaceegg acc 1493

<sup>&</sup>lt;210> 95

<sup>&</sup>lt;211> 177

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapien

<sup>&</sup>lt;400> 95

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Met Asn Ser Gly Lys Arg Arg Leu Pro Trp Arg Leu Arg Ser Gly Val

Pro Ser Pro Pro Gly Leu Leu Ala Pro Ala Pro Ala Pro Cys Ala Pro

Gly Gly His Arg Arg Ala Pro Gly Pro Arg Arg Val Arg Glu Thr Pro 40

Arg Thr Gly Gly Gly Ile Gly Pro Pro Ser Phe Gly Gly Lys Gly

Gly Trp Lys Glu Glu Gly Ser Gly Val Gly Glu Ser Trp Ser Phe Gly

Ile Phe Ser Pro Gly Gln Ala Val Leu Arg Ala Leu Arg Cys Val Ser 85

Lys Cys Trp Glu Asp Ser Ala Gly Lys Gly Leu Arg Thr Arg Pro Ala 105

Gly Thr Gly Val Ala Ala Ser Glu Gly Arg Gly Glu Pro Met Ala Ser 120

Arg Leu Trp Thr Arg Arg Pro Ser Pro Gly Arg Ser Ala Arg Ser Pro 130

Pro Pro Ala Ser Cys Ala Gly Pro Cys Pro Ala Ser Pro Ala Met Val 155 145

Pro His Pro Pro Pro Arg Glu Arg Pro Cys Pro Pro Ile Leu His Phe 165

Pro

<210> 96

<211> 55 <212> PRT <213> Homo sapien

<400> 96

Met Gln Asn Ser Thr Ser Ser Gly Leu Cys Val Asn Val Pro Pro Phe 5

Pro Pro Leu Ser Gly Cys Leu Asn Val Phe Pro Phe His Leu Lys

9.1

30 25 20

Leu Cys Leu Asp Val Leu His Cys His His Leu Phe Leu Arg Lys Arg 40

Cys Val Pro His Pro Asn Pro

<210> 97 <211> 24 <212> PRT <213> Homo sapien

<400> 97

Met Asp His Phe Tyr Leu Leu Ser Asp Thr Tyr Leu Leu Gly Cys Glu 5 . 10

Pro Gln Gly Gly Leu Leu Leu Gly 20

<210> 98

<211> 646

<212> PRT

<213> Homo sapien

<400> 98

Met Glu Pro Ala Ala Gly Phe Leu Ser Pro Arg Pro Phe Gln Arg Ala 5

Ala Ala Ala Pro Ala Pro Pro Ala Gly Pro Gly Pro Pro Pro Ser Ala 25

Leu Arg Gly Pro Glu Leu Glu Met Leu Ala Gly Leu Pro Thr Ser Asp

Pro Gly Arg Leu Ile Thr Asp Pro Arg Ser Gly Arg Thr Tyr Leu Lys

Gly Arg Leu Leu Gly Lys Gly Gly Phe Ala Arg Cys Tyr Glu Ala Thr

Asp Thr Glu Thr Gly Ser Ala Tyr Ala Val Lys Val Ile Pro Gln Ser

Arg Val Ala Lys Pro His Gln Arg Glu Lys Ile Leu Asn Glu Ile Glu 105

Leu His Arg Asp Leu Gln His Arg His Ile Val Arg Phe Ser His His 115 120 125

Phe Glu Asp Ala Asp Asn Ile Tyr Ile Phe Leu Glu Leu Cys Ser Arg · 130 135 140

Lys Ser Leu Ala His Ile Trp Lys Ala Arg His Thr Leu Leu Glu Pro 145 150 155 160

Glu Val Arg Tyr Tyr Leu Arg Gln Ile Leu Ser Gly Leu Lys Tyr Leu 165 170 175

His Gln Arg Gly Ile Leu His Arg Asp Leu Lys Leu Gly Asn Phe Phe 180 185 190

Ile Thr Glu Asn Met Glu Leu Lys Val Gly Asp Phe Gly Leu Ala Ala 195 200 205

Arg Leu Glu Pro Pro Glu Gln Arg Lys Lys Thr Ile Cys Gly Thr Pro 210 215 220

Asn Tyr Val Ala Pro Glu Val Leu Leu Arg Gln Gly His Gly Pro Glu 225 230 235 240

Ala Asp Val Trp Ser Leu Gly Cys Val Met Tyr Thr Leu Leu Cys Gly 245 250 255

Ser Pro Pro Phe Glu Thr Ala Asp Leu Lys Glu Thr Tyr Arg Cys Ile 260 265 270

Lys Gln Val His Tyr Thr Leu Pro Ala Ser Leu Ser Leu Pro Ala Arg 275 280 285

Gln Leu Leu Ala Ala Ile Leu Arg Ala Ser Pro Arg Asp Arg Pro Ser 290 295 300

Ile Asp Gln Ile Leu Arg His Asp Phe Phe Thr Lys Gly Tyr Thr Pro 305 310 315 320

Asp Arg Leu Pro Ile Ser Ser Cys Val Thr Val Pro Asp Leu Thr Pro 325 330 335

Pro Asn Pro Ala Arg Ser Leu Phe Ala Lys Val Thr Lys Ser Leu Phe 340 345 350

Gly Arg Lys Lys Ser Lys Asn His Ala Gln Glu Arg Asp Glu Val

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365

355 360

Ser Gly Leu Val Ser Gly Leu Met Arg Thr Ser Val Gly His Gln Asp 370 375 380

Ala Arg Pro Glu Ala Pro Ala Ala Ser Gly Pro Ala Pro Val Ser Leu 385 390 395 400

Val Glu Thr Ala Pro Glu Asp Ser Ser Pro Arg Gly Thr Leu Ala Ser 405 410 415

Ser Gly Asp Gly Phe Glu Glu Gly Leu Thr Val Ala Thr Val Val Glu 420 425 430

Ser Ala Leu Cys Ala Leu Arg Asn Cys Ile Ala Phe Met Pro Pro Ala 435 440 445

Glu Gln Asn Pro Ala Pro Leu Ala Gln Pro Glu Pro Leu Val Trp Val 450 455 460

Ser Lys Trp Val Asp Tyr Ser Asn Lys Phe Gly Phe Gly Tyr Gln Leu 465 470 475 480

Ser Ser Arg Arg Val Ala Val Leu Phe Asn Asp Gly Thr His Met Ala 485 490 495

Leu Ser Ala Asn Arg Lys Thr Val His Tyr Asn Pro Thr Ser Thr Lys 500 505 510

His Phe Ser Phe Ser Val Gly Ala Val Pro Arg Ala Leu Gln Pro Gln 515 520 525

Leu Gly Ile Leu Arg Tyr Phe Ala Ser Tyr Met Glu Gln His Leu Met 530 535 540

Lys Gly Gly Asp Leu Pro Ser Val Glu Glu Val Glu Val Pro Ala Pro 545 550 555 560

Pro Leu Leu Gln Trp Val Lys Thr Asp Gln Ala Leu Leu Met Leu 565 570 575

Phe Ser Asp Gly Thr Val Gln Val Asn Phe Tyr Gly Asp His Thr Lys 580 585 590

Leu Ile Leu Ser Gly Trp Glu Pro Leu Leu Val Thr Phe Val Ala Arg 595 600 605

Asn Arg Ser Ala Cys Thr Tyr Leu Ala Ser His Leu Arg Gln Leu Gly 615

Cys Ser Pro Asp Leu Arg Gln Arg Leu Arg Tyr Ala Leu Arg Leu Leu 635 630

Arg Asp Arg Ser Pro Ala

<210> 99

<211> 99

<212> PRT <213> Homo sapien

<400> 99

Met Leu Thr Ser Pro Ser Thr Tyr Val Ile Gln Glu Asn Gly Ser Leu

Val Glu Ile Arg Asn Ile Leu Gly Glu Lys Tyr Ile Arg Arg Val Arg 25

Met Arg Pro Gly Val Ala Cys Ser Val Ser Gln Ala Gln Lys Asp Glu 35

Leu Ile Leu Glu Gly Asn Asp Ile Glu Leu Val Ser Asn Ser Ala Cys 50

Phe Gly Cys Gln Gln Met Pro Gln Ser Val Lys Asn Lys Asp Ile Arg 70

Lys Phe Leu Asp Gly Ile Tyr Val Ser Glu Lys Gly Thr Val Gln Gln

Ala Asp Glu

<210> 100

<211> 220

<212> PRT

<213> Homo sapien

<400> 100

Met Lys Thr Ile Leu Ser Asn Gln Thr Val Asp Ile Pro Glu Asn Gly 5

Met Arg Leu Asp Val Phe Tyr Leu His Leu Tyr Cys Thr Phe Gln Ala

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20

Leu Cys Gly Leu Thr Ser Val Phe Ser Leu Leu Val Asp Ile Thr Leu

25

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30

Lys Gly Arg Thr Val Ile Val Lys Gly Pro Arg Gly Thr Leu Arg Arg

Asp Phe Asn His Ile Asn Val Glu Leu Ser Leu Leu Gly Lys Lys

Lys Arg Leu Arg Val Asp Lys Trp Trp Gly Asn Arg Lys Glu Leu Ala 90

Thr Val Arg Thr Ile Cys Ser His Val Gln Asn Met Ile Lys Gly Val

Thr Leu Gly Phe Arg Tyr Lys Met Arg Ser Val Tyr Ala His Phe Pro 120

Ile Asn Val Val Ile Gln Glu Asn Gly Ser Leu Val Glu Ile Arg Asn 135 130

Phe Leu Gly Glu Lys Tyr Ile Arg Arg Val Arg Met Arg Pro Gly Val 155 150

Ala Cys Ser Val Ser Gln Ala Gln Lys Asp Glu Leu Ile Leu Glu Gly 165 170

Asn Asp Ile Glu Leu Val Ser Asn Ser Ala Ala Leu Ile Gln Gln Ala

Thr Thr Val Lys Asn Lys Asp Ile Arg Lys Phe Leu Asp Gly Ile Tyr

Val Ser Glu Lys Gly Thr Val Gln Gln Ala Asp Glu 215 210

<210> 101

<211> 47

<212> PRT

<213> Homo sapien

<400> 101

Met Arg Trp His Thr Tyr Leu Cys Cys Leu Lys Val Thr Ile Met Leu

96

Pro Tyr Gln Ala Glu Asn Val Thr Thr Ile Trp Arg Phe Arg Arg Val

Phe Leu Ser Glu Ser Val Met Asn Thr Leu Val Gly Trp Ile Gln 40

<210> 102

<211> 51

. <212> PRT

<213> Homo sapien

<400> 102

Met Ser Ser His Lys Thr Phe Arg Ile Lys Arg Phe Leu Ala Lys Lys

Gln Lys Gln Asn Arg Pro Ile Pro Gln Trp Ile Arg Met Lys Thr Gly

Asn Lys Ile Arg Tyr Asn Ser Lys Arg Arg His Trp Arg Arg Thr Lys 40

Leu Gly Leu 50

<210> 103 <211> 53 <212> PRT <213> Homo sapien

<400> 103

Met Glu Arg Val Leu Glu Lys Gln Glu Lys Lys Ser Cys Leu Lys Pro

His Val Tyr Cys Arg His Arg Arg Glu Trp Arg His Leu Ser Ile Leu 20 25

Phe Ser Ile Ser Thr Ala Pro Gln Asn Thr Tyr Ile Leu Phe Phe

Phe Ser Glu Met Ser 50

<210> 104 <211> 131

<212> PRT

<213> Homo sapien

<400> 104

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Met Arg Val Ser Glu Arg Ala Leu Lys Asn Val Ala Cys Gln Gln His 1 5 10 15

Met Asp Ser Leu Phe Arg Val Cys Ile Tyr Pro Ala Asp Thr Pro Ile 20 25 30

Pro Pro Ser Leu Pro Pro Arg Ala Ser Asp Phe Leu Phe His Pro Ala 35 40 45

Ala Tyr Tyr Trp Gln Gly Met Ala Gly Val Asn Leu Gly Ser Val Tyr 50 55 60

His Gln Gly Lys Leu Pro Ser Leu Leu Gln Ser Leu Trp Lys Gly Thr 65 70 75 80

Phe Phe Arg Val Gln His Val Pro Met His Ser Gln Val Pro Lys Val 85 90 95

Thr Tyr Thr Tyr Ile Val Asn Ile Val Pro Thr Ala Leu Gln Thr Phe
100 105 110

Ile Trp Pro Leu Ala Val His Thr Ser Gln Pro Ile His Val Phe Met
115 120 125

Met Met Phe 130

<210> 105

<211> 117

<212> PRT

<213> Homo sapien

<400> 105

Met Ser Ser Phe Gln Gly Phe Ile Phe Gly Gly Lys Lys Ile Pro Gln 1 5 10 15

Asp Ala Gly Cys Pro Ala Ser His Asn Gly Tyr Ala Pro Ile Glu Thr 20 25 30

Ser Ser Gly Arg Val Thr Lys Leu Lys Arg Lys Gln Phe Gln Ala Glu 35 40 45

Gly His Lys Leu Arg Ala Glu Ser Leu Leu Leu Thr Ala Ile Gln Ala 50 55 60

Gln Gly Leu Cys Gly Ala Gly Phe Leu Lys Ala Gly Leu Tyr Leu Gly

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80 75 65 70

Arg Arg Glu Arg Thr Arg Gly Leu Asp Ala Gly Trp Arg Phe Cys Asp

Leu Leu Cys Tyr Lys Phe Lys Asn Lys Thr Cys Trp Ile Arg Ser Phe 100 105

Ser Tyr Leu Leu Lys 115

<210> 106

<211> 93 <212> PRT <213> Homo sapien

<400> 106

Met Pro Gly Val Thr Val Lys Asp Val Asn Gln Glu Phe Val Arg

Ala Leu Ala Ala Phe Leu Lys Lys Ser Gly Glu Ala Glu Ser Pro Arg 25

Met Gly Gly Ile Pro Phe Lys Leu Ala Lys Ala Gln Arg Ser Leu Leu

Pro Thr Met Arg Thr Gly Ser Thr Arg Gly Ala Ala Phe Gln Gln Arg

Arg Ala Thr Cys Tyr Leu Pro Gly Val Gly Ala Gly Gly Trp Ala Ser

Ile Glu Pro Lys Asp Ser Ile Gly Gly Glu Arg Ser Glu

<210> 107 <211> 148 <212> PRT

<213> Homo sapien

<400> 107

Met Leu Leu Val Gly Ser Cys His Leu Ser Gly Asp Ser Val Gln Ile

Ser Leu Ser Leu Arg Cys Gln Phe Ala Ala Ala Ile Leu Val Leu Phe . 25

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His His Phe Gln Pro Leu Gln Gly Leu Glu Asp Pro Ala Gly His Thr 35 40 45

Leu Gly Ala Ser Ala Glu Val Ala Gly His Asp Ala Val Ser Leu Thr 50 55 60

Ser Pro Ile Asp Leu Gly His Gly Ala Asn Pro Ser Ala Thr Pro Glu 65 70 75 80

Val Gln Val Pro Arg Cys Gly Ser Ser Ser Arg Val Glu Pro Val Leu 85 90 95

Ile Val Gly Ser Lys Leu Phe Val Leu Gly Gln Leu Asp Gly Ile His
100 105 110

Pro Phe Gly Asp Phe Gln Leu Pro Gly Leu Phe Glu Glu Gly Cys Gln 115 120 125

Ser Ser Asp Glu Leu Leu Val His Val Phe Tyr Ser Asn Ser Arg 130 135 140

His Arg Ala Ala 145

<210> 108

<211> 172

<212> PRT

<213> Homo sapien

<400> 108

Met Val Cys Gly Gly Phe Ala Cys Ser Ser Leu Arg Val Val Gly Val 1 5 10 15

Val Ile Ala Val Gly Ile Phe Leu Phe Leu Ile Ala Leu Val Gly Leu 20 25 30

Ile Gly Ala Val Lys His His Gln Val Leu Leu Phe Phe Tyr Met Ile 35 40 45

Ile Leu Leu Leu Val Phe Ile Val Gln Phe Ser Val Ser Cys Ala Cys 50 55 60

Leu Ala Leu Asn Gln Glu Gln Gln Gly Gln Leu Leu Glu Val Gly Trp 65 70 75 80

Asn Asn Thr Ala Ser Ala Arg Asn Asp Ile Gln Arg Asn Leu Asn Cys 85 90 95

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Cys Gly Phe Arg Ser Val Asn Pro Asn Asp Thr Cys Leu Ala Ser Cys 105

Val Lys Ser Asp His Ser Cys Ser Pro Cys Ala Pro Ile Ile Gly Glu 120 115

Tyr Ala Gly Glu Val Leu Arg Phe Val Gly Gly Ile Gly Leu Phe Phe 130

Ser Phe Thr Glu Ile Leu Gly Val Trp Leu Thr Tyr Arg Tyr Arg Asn 155 150 145

Gln Lys Asp Pro Arg Ala Asn Pro Ser Ala Phe Leu 165

<210> 109

<211> 55

<212> PRT <213> Homo sapien

<400> 109

Met Asn Phe Asp Tyr Ser Val Asn Tyr Trp Asn Val Ser Ser Phe Asn 5

Phe Lys Asn Asn Tyr Phe Thr Ser Ser Asp Trp Gly Phe Pro Glu Ile

Cys Glu Glu Gln Arg Arg Pro Pro Ala Thr Gln His His Asp Gly 35

Ala Leu Thr Gly Ser Glu Ser

<210> 110 <211> 125 <212> PRT

<213> Homo sapien

Met Gln Ala Leu Pro Gln Val Glu Lys Arg Arg Leu Arg Leu Pro Arg 1 5

Glu Val Gln Cys Pro Ala Leu Leu Arg Arg Met Leu Leu Ile Pro Leu 20

Trp Lys Ile Pro Ala Pro Thr Thr Lys Ser Cys Arg Glu Thr Phe

35 40 45

Leu Lys Trp Leu Ser Val Ser Ala Ala Glu Arg Thr Thr Gly Ser Trp 50 55 60

Thr Ser Phe Gln Pro Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro 65 70 75 80

Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg 85 90 95

Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser 100 105 110

Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu 115 120 125

<210> 111

<211> 1256

<212> PRT

<213> Homo sapien

<400> 111

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr
1 5 10 15

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 20 25 30

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser 35 40 45

Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser His 50 55 60

Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr Leu 65 70 75 80

Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln 85 90 95

Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu Gly Ser Thr Thr 100 105 110

Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Lys Pro Ala Pro 115 120 125

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 130 135 140

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 145 150 155 160

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 165 170 175

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Pro Pro Gly Ser Thr Ala 180 185 190

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 195 200 205

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 210 215 220

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 225 230 235 240

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 245 250 255

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 260 265 270

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 275 280 285

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 290 295 300

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 305 310 315 320

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 325 330 335

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 340 345 350

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 355 360 365

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 370

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 390

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 410 405

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 425 420

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 445 440 435

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 450 455

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 485 490

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 505

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 550

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 570 575 565

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 580 585

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 600 605 595

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr

104

610 615 620

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 625 630 635

- Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His
  645 650 655
- Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 660 665 670
- Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 675 680 685
- Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 690 695 700
- Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 705 710 715 720
- Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His
  725 730 735
- Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 740 745 750
- Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 755 760 765
- Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 770 775 780
- Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 785 790 795 800
- Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 805 810 815
- Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 820 825 830
- Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 835 840 845
- Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 850 855 860

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 865 870 875 880

- Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Val His 885 890 895
- Gly Val Thr Ser Ala Pro Asp Ser Arg Ser Gly Ser Gly Phe Leu Pro 900 905 910
- Pro Pro Ala Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala 915 920 925
- Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp 930 935 940
- Asn Arg Pro Ala Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr 945 950 955 960
- Ser Ala Ser Gly Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn 965 970 975
- Gly Thr Ser Ala Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro 980 985 990
- Phe Ser Ile Pro Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser
- His Ser Thr Lys Thr Asp Ala Ser Ser Thr His His Ser Thr Val 1010 1015 1020
- Pro Pro Leu Thr Ser Ser Asn His Ser Thr Ser Pro Gln Leu Ser 1025 1030 1035
- Thr Gly Val Ser Phe Phe Phe Leu Ser Phe His Ile Ser Asn Leu 1040 1045 1050
- Gln Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp Tyr Tyr Gln 1055 1060 1065
- Glu Leu Gln Arg Asp Ile Ser Glu Met Phe Leu Gln Ile Tyr Lys 1070 1075 1080
- Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly 1085 1090 1095

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Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile 1105 1100

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Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu 1120

Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser 1135

Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val Pro 1150

Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala

Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg 1180 1175

Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr 1195 1200 1190

Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg 1210 1215 1205

Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val 1230 1220 1225

Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala 1240 1245 1235

Val Ala Ala Thr Ser Ala Asn Leu 1255 1250

<210> 112

<211> 728

<212> PRT <213> Homo sapien

<400> 112

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 20 25 30

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser

107

...

45 40 35 Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser His 55 Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Lys Pro Ala Pro 120 Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 135 Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Pro Ala His Gly Val Thr 150 155 Ser Ala Pro Asp Thr Arg Pro Pro Pro Gly Ser Thr Ala Pro Pro Ala 165 170 His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 185 180 Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 200 195 Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 210 215 Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 275 280 285

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- Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 290 295 300
- Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 305 310 315 320
- Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 325 330 335
- Met Val Ser Ile Gly Leu Ser Phe Pro Ser Ser Pro Glu Ala Ala Ile 340 345 350
- Arg Thr Val His Thr Leu Cys Ile Lys Pro Glu Ser Phe Pro Ser His 355 360 365
- Pro Ser Phe Cys Arg Phe Ile Asn Lys Gly Val Phe Trp Ala Ser Pro 370 375 380
- Ile Leu Ser Ser Gly Thr Val Leu Gly Val Asp Pro Val Trp Trp Leu 385 390 395 400
- Glu Gly Trp Val Val Val Met Thr Val Gly Gly Thr Gly Arg Thr Tyr 405 410 415
- Gly Trp Gly Lys Ser Arg Glu Pro Glu Leu Gly Pro Val Ala Glu Val 420 425 430
- Pro Ile Phe Pro Val Thr Arg Pro Gly Ser Val Val Val Gln Leu Thr 435 440 445
- Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln 450 460
- Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile 465 470 475 480
- Ser Asp Val Ser Gly Glu Ala Thr Ser Leu Ala Ala Ala Gln His His 485 490 495
- Ala Gly Ala Leu Ser Phe Gln Cys Leu Gly Pro Arg Ser Phe Leu Ser 500 505 510
- Ala Gly Ser Gly Arg Gly Ala Ser Ser Gly Arg Leu Pro Cys Pro Leu 515 520 525

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Leu Phe Leu Leu Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser 530 535 540

Gly Ala Gly Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys 545 550 555 560

Val Leu Val Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Val Ser Ala 565 570 575

Val Pro Gly Pro Asp Gln Ser Pro Pro Val Glu Gly Ser Ser Met Ala 580 585 590

Cys His Asn Leu Leu Ser Pro Gln Ala Val Cys Gln Cys Arg Arg Lys 595 600 605

Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro 610 615 620

Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro 625 630 635 640

Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Arg Leu Ala Pro Gln 645 650 655

Ala Arg Gly Ser Arg Gly Phe Gly Trp Ala Arg Ile Leu Lys Gly Val 660 665 670

Leu Gly Lys Pro Lys Glu Leu Gly Arg Gly Glu Lys Trp Arg Glu Val 675 680 685

Ser Arg Gly Gly Pro Gly Lys Asp Glu Gly Gln Arg Ser Glu Glu Phe 690 695 700

Trp Gly Thr Gly Leu Gly Gly Asp Tyr Gly Arg Lys Gly Pro Ser Lys 705 710 715 720

Gly Ser Gly Pro Thr Ala Arg Ile 725

<210> 113

<211> 524

<212> PRT

<213> Homo sapien

<400> 113

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr

110

1 5 10 15

Val Leu Thr Ala Thr Thr Ala Pro Thr Pro Ala Thr Val Val Thr Gly 20 25 30

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 35 40 45

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50 55 60

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65 70 75 80

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85 90 95

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 100 105 110

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 150 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala 165 170 175

Pro Gly Ser Thr Ala Pro Ala Ala His Gly Val Thr Ser Ala Pro Asp 180 185 190

Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr 195 200 205

Ser Ala Pro Asp Asn Arg Pro Ala Leu Gly Ser Thr Ala Pro Pro Val 210 215 220

His Asn Val Thr Ser Ala Ser Gly Ser Ala Ser Gly Ser Ala Ser Thr 225 230 235 240

Leu Val His Asn Gly Thr Ser Ala Arg Ala Thr Thr Thr Pro Ala Ser 245 250 255

- Lys Ser Thr Pro Phe Ser Ile Pro Ser His His Ser Asp Thr Pro Thr
- Thr Leu Ala Ser His Ser Thr Lys Thr Asp Ala Ser Ser Thr His His 280
- Ser Thr Val Pro Pro Leu Thr Ser Ser Asn His Ser Thr Ser Pro Gln 295 290
- Leu Ser Thr Gly Val Ser Phe Phe Phe Leu Ser Phe His Ile Ser Asn 305 310
- Leu Gln Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp Tyr Tyr Gln 330
- Glu Leu Gln Arg Asp Ile Ser Glu Met Phe Leu Gln Ile Tyr Lys Gln 340 345
- Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val
- Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His 375 370
- Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Arg
- Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Pro 410
- Phe Ser Ala Gln Ser Gly Ala Gly Val Pro Gly Trp Gly Ile Ala Leu 425
- Leu Val Leu Val Cys Val Leu Val Ala Leu Ala Ile Val Tyr Leu Ile 440
- Ala Leu Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp 455 450
- Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro Thr 470 475 480 465
- Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser 490 485

112

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Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr 505

Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu

<210> 114

<211> 515 <212> PRT

<213> Homo sapien

<400> 114

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 25 20

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser

Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser His 60

Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr Leu

Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln

Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu Gly Ser Thr Thr 105

Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Lys Pro Ala Pro

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 150 145

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Ala Ala His 165 170

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala

113

185 190 180 Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala Leu 200 Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala Arg 235 Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro Ser 245 His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys Thr 265 270 Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser Ser 280 Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 295 290 Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu Asp 305 310 315 320 Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu Met 330 325 Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr 370 Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val 410 405 Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala 425

Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg

Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His

Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro 465 470

Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn 490 485

Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser 505 510

Ala Asn Leu 515

<210> 115

<211> 109 <212> PRT <213> Homo sapien

<400> 115

Met Leu Glu Arg Arg Pro Pro Ala Val Arg Arg Pro Gly Leu Thr Ala 5 10

Pro Ala Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro

Gly Ser Thr Ala Pro Ala Ala His Gly Val Thr Ser Ala Pro Asp Thr 35

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Phe

Val Pro Arg Thr Ser Gly Arg Arg Leu Ala Leu Phe Leu Val Tyr Val 70

Phe Arg Val Glu Asp Val Val Gln Thr Arg Leu Asp Thr Leu Arg Ile 90

Ala Lys Tyr Ile Asp Gly Ser Tyr Ala Val Ser Val Cys 105

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<210> 116

<211> 174

<212> PRT

<213> Homo sapien

<220>

<221> MISC\_FEATURE

<222> (167)..(167) <223> X= any amino acid

<400> 116

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Ala Thr Thr Ala Pro Thr Pro Ala Thr Val Val Thr Gly 25

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50 55 60

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 70

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 105 100

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Arg Pro 140 135

Ser Cys Gly Ser Gly Leu Gly Thr Ala Cys Val Pro Gly Leu Gln Leu 145

Leu Leu Val Gly Ala His Xaa Thr Gln Leu Leu Thr Tyr Asp 165

<210> 117 <211> 475 <212> PRT

<213> Homo sapien

<400> 117

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Leu Thr 1 5 10 15

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly
20 25 30

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser 35 40 45

Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser His 50 55 60

Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr Leu 65 70 75 80

Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln 85 90 95

Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu Gly Ser Thr Thr 100 105 110

Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Lys Pro Ala Pro 115 120 125

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 130 135 140

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 145 150 155 160

Ala Pro Asp Asn Arg Pro Ala Leu Gly Ser Thr Ala Pro Pro Val His 165 170 175

Asn Val Thr Ser Ala Ser Gly Ser Ala Ser Gly Ser Ala Ser Thr Leu 180 185 190

Val His Asn Gly Thr Ser Ala Arg Ala Thr Thr Thr Pro Ala Ser Lys 195 200 205

Ser Thr Pro Phe Ser Ile Pro Ser His His Ser Asp Thr Pro Thr Thr 210 215 220

Leu Ala Ser His Ser Thr Lys Thr Asp Ala Ser Ser Thr His His Ser 225 230 240

Thr Val Pro Pro Leu Thr Ser Ser Asn His Ser Thr Ser Pro Gln Leu 245 250 255

Ser Thr Gly Val Ser Phe Phe Phe Leu Ser Phe His Ile Ser Asn Leu 260 265 270

Gln Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp Tyr Tyr Gln Glu 275 280 285

Leu Gln Arg Asp Ile Ser Glu Met Phe Leu Gln Ile Tyr Lys Gln Gly 290 295 300

Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val Val 305 310 315 320

Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp 325 330 335

Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Arg Tyr 340 345 350

Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Pro Phe 355 360 365

Ser Ala Gln Ser Gly Ala Gly Val Pro Gly Trp Gly Ile Ala Leu Leu 370 375 380

Val Leu Val Cys Val Leu Val Ala Leu Ala Ile Val Tyr Leu Ile Ala 385 390 395 400

Leu Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile 405 410 415

Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr 420 425 430

His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro 435 440 445

Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr 450 455 460

Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu 465 470 475

<210> 118

<211> 231 <212> PRT <213> Homo sapien

<400> 118

Met Cys Pro Leu Ala Val Pro Ile Val Ala Pro Met Arg Arg Phe Leu

Gln Val Met Val Ala Ala Ala Ser Leu Thr Gln Thr Gln Gln Trp Gln

Pro Leu Leu Pro Thr Cys Arg Gly Thr Ser Pro Ala Glu Leu Ser Gly 40 35

Gln Pro Val Pro Phe His Ser Thr Gln Val Leu Gln Gly Gln Ser Pro 55

Cys Thr Leu Phe Gly Leu Val Ser Trp Glu Phe Arg Trp Ala Ala His

Ser Leu Leu Gln Arg Pro His Asp Tyr Phe Arg Lys Phe Glu Pro His

Leu Tyr Ser Leu Asp Ser Asn Ser Asp Asp Val Asp Ser Leu Thr Asp

Glu Glu Ile Leu Ser Lys Tyr Gln Leu Gly Met Leu His Phe Ser Thr 120

Gln Tyr Asp Leu Leu His Asn His Leu Thr Val Arg Val Ile Glu Ala 135

Arg Asp Leu Pro Pro Pro Ile Ser His Asp Gly Ser Arg Gln Asp Met 150 155

Ala His Ser Asn Pro Tyr Val Lys Ile Cys Leu Leu Pro Asp Gln Lys 165 170

Asn Ser Lys Gln Thr Gly Val Lys Arg Lys Thr Gln Lys Pro Val Phe 180

Glu Glu Arg Tyr Thr Phe Glu Ile Pro Phe Leu Glu Ala Gln Arg Arg 195 200 205

Thr Leu Leu Thr Val Val Asp Phe Asp Lys Phe Ser Arg His Cys

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119

220 210 215

Val Ile Gly Lys Val Ser Val

<210> 119

<211> 107 <212> PRT <213> Homo sapien

<400> 119

Met Val Ala Ala Ser Leu Thr Gln Thr Gln Gln Trp Gln Pro Leu

Leu Pro Thr Cys Arg Gly Thr Ser Pro Ala Glu Leu Ser Gly Gln Pro

Val Pro Phe His Ser Thr Gln Val Leu Gln Gly Gln Ser Pro Cys Thr 40 35

Leu Phe Gly Leu Val Ser Trp Glu Phe Arg Trp Ala Ala His Ser Leu

Leu Gln Arg Pro His Gln Phe Leu Gly His Phe Ser Val Cys Gly Ser

Ser Cys Gly Pro Leu Arg Ala His Ala Trp Glu Val Leu Trp Trp Gly

Leu Pro Gly Gly Leu Ala Gln Arg Ala Leu Arg 1,95

<210> 120 <211> 484

<212> PRT

<213> Homo sapien

<400> 120

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Leu Thr

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 25

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 40

120

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50 55 60

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65 70 75 80

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85 90 95

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val
100 105 110

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr. Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly
180 185 190

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200 205

Arg Ala Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 275 280 285

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu

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300 295 290

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn

Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe 330

Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln

Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val 360

Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly

Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val 390

Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg 415

Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr 425 420

His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val 440 435

Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly 460 455 450

Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr 470 465

Ser Ala Asn Leu

<210> 121

<211> 463 <212> PRT <213> Homo sapien

<400> 121

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Gly Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser 20 25 30

Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val 35 40 45

Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln 50 55 60

Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala 65 70 75 80

Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu 85 90 95

Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn 100 105 110

Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 115 120 125

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 130 135 140

Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala Leu Gly Ser Thr Ala 145 150 155 160

Pro Pro Val His Asn Val Thr Ser Ala Ser Gly Ser Ala Ser Gly Ser 165 170 175

Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala Arg Ala Thr Thr 180 185 190

Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro Ser His His Ser Asp 195 200 205

Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys Thr Asp Ala Ser Ser 210 215 220

Thr His His Ser Thr Val Pro Pro Leu Thr Ser Ser Asn His Ser Thr 225 230 235 240

Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe Phe Leu Ser Phe His 245 250 255

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Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp 265 260

Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu Met Phe Leu Gln Ile 280

Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro 300

Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile 310 305

Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala 325

Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val 345

Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val Pro Gly Trp Gly 355

Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala Leu Ala Ile Val 375

Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly 385 390

Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu

Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr 425

Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser

Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu

<210> 122 <211> 524 <212> PRT

<213> Homo sapien

<400> 122

Met Gly Arg Glu Lys Glu Ala Ala Ala Gly Lys Glu Ala Ala Asn Pro

Gly Val Thr Glu Ala Ala His Ser Pro Val Leu Leu Val Leu Phe Leu 20 25 30

Trp Trp Pro Glu Leu Ile Phe Ser Ser Cys Ser Tyr Phe Ser Phe Ile 35 40 45

Lys Thr Gln Pro Tyr Asp Phe Asn Phe Phe Thr Ala Thr Thr Ala Pro 50 55 60

Lys Pro Ala Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro 65 70 75 80

Gly Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser 85 90 95

Ser Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser 100 105 110

His Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr

Leu Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly
130 135 140

Gln Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu Gly Ser Thr 145 150 155 160

Thr Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Lys Pro Ala 165 170 175

Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp 180 185 190

Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr 195 200 205

Ser Ala Pro Asp Asn Arg Pro Ala Leu Gly Ser Thr Ala Pro Pro Val 210 215 220

His Asn Val Thr Ser Ala Ser Gly Ser Ala Ser Gly Ser Ala Ser Thr 225 230 235 240

Leu Val His Asn Gly Thr Ser Ala Arg Ala Thr Thr Thr Pro Ala Ser 245 250 255

125

- Lys Ser Thr Pro Phe Ser Ile Pro Ser His His Ser Asp Thr Pro Thr 260 265 270
- Thr Leu Ala Ser His Ser Thr Lys Thr Asp Ala Ser Ser Thr His His 275 280 285
- Ser Thr Val Pro Pro Leu Thr Ser Ser Asn His Ser Thr Ser Pro Gln 290 295 300
- Leu Ser Thr Gly Val Ser Phe Phe Phe Leu Ser Phe His Ile Ser Asn 305 310 315 320
- Leu Gln Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp Tyr Tyr Gln 325. 330 335
- Glu Leu Gln Arg Asp Ile Ser Glu Met Phe Leu Gln Ile Tyr Lys Gln 340 . 345 350
- Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val 355 360 365
- Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His 370 375 380
- Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Arg 385 390 395 400
- Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Pro 405 410 415
- Phe Ser Ala Gln Ser Gly Ala Gly Val Pro Gly Trp Gly Ile Ala Leu 420 425 430
- Leu Val Leu Val Cys Val Leu Val Ala Leu Ala Ile Val Tyr Leu Ile 435 440 445
- Ala Leu Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp 450 460
- Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro Thr 465 470 475 480
- Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser 485 490 495

126

Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr 505 500

Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu

<210> 123

<211> 435 <212> PRT <213> Homo sapien

<400> 123

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser

Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser His 55

Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr Leu 75 70

Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln 85

Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu Gly Ser Thr Thr 105 100

Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Arg Pro Ala Leu 115 120

Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly Ser 135

Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala Arg 145 150 155

Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro Ser 170

His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys Thr 185 190 180

Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser Ser . 195 200 205

Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 210 215 220

Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu Asp 225 230 235 240

Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu Met 245 250 255

Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile 260 265 270

Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg 275 280 285

Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr 290 295 300

Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser 305 310 315 320

Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val 325 330 335

Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala 340 345 350

Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg 355 360 365

Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His 370 375 380

Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro 385 390 395 400

Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn 405 410 415

Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser
420 430

Ala Asn Leu 435

<210> 124

<211> 273

<212> PRT

<213> Homo sapien

<400> 124

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Leu Thr 1 5 10 15

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 20 25 30

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser 35 40 45

Thr Glu Lys Asn Ala Leu Ser Thr Gly Val Ser Phe Phe Leu Ser 50 55 60

Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu Asp Pro Ser 65 70 75 80

Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu Met Phe Leu 85 90 95

Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe 100 105 110

Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly 115 120 125

Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr 130 135 140

Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser 145 150 155 160

Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val Pro Gly 165 170 175

Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala Leu Ala 180 185 190

Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg Lys Asn

129

195 200 205

Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met 210 215 220

Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser 225 230 235 240

Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly
245 250 255

Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn 260 265 270

Leu

<210> 125

<211> 350

<212> PRT

<213> Homo sapien

<400> 125

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 1 5 10 15

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20 25 30

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 35 40 45

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50 55 60

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65 70 75 80

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85 90 95

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val
100 105 110

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly
180 185 190

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200 205

Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 275 280 285

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 290 295 300

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 305 310 315 320

Ile Lys Phe Arg Tyr Ser Ser Gly Cys Gly Pro Ser Val Val Gly 325 330 335

Gly Trp Val Val Val Met Thr Val Gly Arg Asp Trp Cys Thr 340 345 350

<210> 126

<211> 316

<212> PRT

<213> Homo sapien

<400> 126

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 1 5 10 15

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly
20 25 30

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50 55 60

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65 70 75 80

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85 90 95

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly
180 185 190

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala
195 200 205

Arg Ala Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 230 235 240

132

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 280

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 295 290

Met Val Ser Ile Gly Leu Ser Phe Pro Met Leu Pro 310

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<210> 127 <211> 230 <212> PRT <213> Homo sapien

<400> 127

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 20 25

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser 35

Thr Glu Lys Asn Ala Ile Pro Ala Pro Thr Thr Lys Ser Cys Arg 50

Glu Thr Phe Leu Lys Trp Pro Gly Ser Val Val Val Gln Leu Thr Leu 70 65

Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe

Asn Gln Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser 105 100

Asp Val Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly 125 120

133

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Ala Gly Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val 135 130

Leu Val Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln

Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp 170

Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg 180

Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser 195 200

Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala 220 215

Ala Thr Ser Ala Asn Leu 225

<210> 128

WO 03/106648

<211> 614 <212> PRT <213> Homo sapien

<400> 128

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20 25

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 35

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 70

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 105 110

- Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115
- Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 140
- His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 150
- Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala
- Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 185
- Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 200 195
- Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 215
- Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 230
- Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 250
- Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 265
- Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu
- Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 295
- Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 315 320 310
- Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe 325
- Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln 345 340

Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val

Ser Val Leu Leu Ile Gly Gly Glu Arg Arg Tyr Arg Ala Met Val 370 375 380

Ser Ala Thr Gly Ile Ser Leu Gly Ala Met Ala Gly Lys Gly Gly 385 390 395 400

Val Ser Glu Trp Trp Leu Gly Ile Glu Asn Gly Val Leu Leu Leu Ala 405 410 415

Gly Val Val Val Ala Leu Ala Glu Val Pro Leu Cys Thr Arg Val Glu
420 425 430

Ala Glu Pro Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu
435
440
445

Thr Ser Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser 450 455 460

Phe Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser 465 470 475 480

Leu Glu Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile 485 490 495

Ser Glu Asp Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly 500 505 510

Ala Gly Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val 515 520 525

Leu Val Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln 530 535 540

Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp 545 550 555 560

Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg
565 570 575

Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser 580 585 590

136

Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala 595 600 605

Ala Thr Ser Ala Asn Leu 610

<210> 129

<211> 372

<212> PRT

<213> Homo sapien

<400> 129

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 1 5 10 15

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20 25 30

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 35 . 40 45

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50 55 60

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65 70 75 80

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85 90 95

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 100 105 110

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 180 185 190

137

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200

Arg Ala Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 235 230

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 265 270 260

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 280 275

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 295

Met Trp Gly Ala Arg Leu Gly His Arg Ala Ala Gly Ala Gly Leu Cys 310

Ser Gly Cys Ala Gly His Cys Leu Ser His Cys Leu Gly Cys Leu Ser 325

Val Pro Pro Lys Glu Leu Arg Ala Ala Gly His Leu Ser Ser Pro Gly

Tyr Leu Pro Ser Tyr Glu Arg Val Pro His Leu Pro His Pro Trp Ala 360 355

Leu Cys Ala Pro 370

<210> 130

<211> 256 <212> PRT <213> Homo sapien

<400> 130

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 10

138

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 20 25 30

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser 35 40 45

Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser His 50 55 60

Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr Leu 65 70 75 80

Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln 85 90 95

Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu Gly Ser Thr Thr

Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Arg Pro Ala Leu 115 120 125

Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly Ser 130 135 140

Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala Arg 145 150 155 160

Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro Ser 165 170 175

His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys Thr 180 185 190

Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser Ser 195 200 205

Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 210 215 220

Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu Asp 225 230 235 240

Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu Met 245 250 255

<210> 131

<211> 492 <212> PRT <213> Homo sapien

<400> 131

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 25

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 35

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 55 50

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 70

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 105

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr.

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 170

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 185

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195

Arg Ala Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 215 210

140

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 275 280 285

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 290 295 300

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 305 310 315 320

Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe 325 330 335

Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln 340 345 350

Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val 355 360 365

Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly 370 375 380

Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val 385 390 395 400

Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg 405 410 415

Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr 420 425 430

His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val 435 440 445

Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Arg Leu Gly 450 455 460

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Pro Thr Gly Gln Gly Lys Gln Arg Val Trp Leu Gly Lys Asp Ser Glu 470

Gly Gly Thr Trp Lys Thr Gln Arg Ala Trp Lys Arg 485

<210> 132

<211> 483

<212> PRT <213> Homo sapien

<400> 132

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 25

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 55

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 75 70 65

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 90

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 105

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 185

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200 205

Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 275 280 285

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 290 295 300

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 305 310 315 320

Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe 325 330 335

Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln 340 345 350

Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val 355 360 365

Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly 370 375 380

Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val 385 390 . 395 400

Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg 405 410 415

Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr 420 425 430

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143

PCT/US03/18934

His Pro Met Ser Glu Trp Arg Val Tyr Glu Glu Lys Lys Lys Glu Val 440 435

Pro Ala Val Pro Glu Thr Leu Lys Lys Lys Arg Arg Asn Phe Ala Glu

Leu Lys Ile Lys Arg Leu Arg Lys Lys Phe Ala Lys Arg Cys Phe Glu 475 470

Arg Gln Gly

<210> 133

<211> 150

<212> PRT <213> Homo sapien

<400> 133

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 25

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser

Thr Glu Lys Asn Ala Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp

Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu Met Ala Val Cys Gln

Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp

Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg 105

Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser 120 115

Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala 135 140 130

Ala Thr Ser Ala Asn Leu

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144

145 150

<210> 134

<211> 168

<212> PRT

<213> Homo sapien

<400> 134

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 10 5

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 20 25

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser 35

Thr Glu Lys Asn Ala Leu Ser Thr Gly Val Ser Phe Phe Leu Ser 55 50

Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu Asp Pro Ser

Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu Met Ala Val 90

Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala 100

Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His

Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys 135

Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala

Val Ala Ala Thr Ser Ala Asn Leu

<210> 135 <211> 79

<212> PRT

<213> Homo sapien

<400> 135

145

Ser Pro Glu Trp Leu Thr Leu Ile Ser Ser Pro Gly Lys Asn Tyr Gly 10

Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu 25

Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr 35 40

Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser 55 50

Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu 70

<210> 136

<211> 398 <212> PRT <213> Homo sapien

<400> 136

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20 25

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 35

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 70

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 105 100

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 140 135 130

146

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 180 185 190

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200 205

Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 275 280 285

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 290 295 300

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 305 310 315 320

Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe 325 330 335

Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln 340 345 350

Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val 355 360 365

Ser Ala Glu Val Pro Phe His Ile Met Leu Thr Asn Met Gly Thr Met 370 375 380

147

Glu Tyr His Asn Val Gly Ala Ile Arg Phe Arg His Asn Tyr 385 390 395

<210> 137

<211> 36

<212> PRT

<213> Homo sapien

<400> 137

Gly Arg Leu Leu Leu Leu Leu Glu Phe Lys Leu Leu Thr Met Tyr 1 5 10 15

Gly Leu Met Pro Gly Lys Cys Cys Gly Gly Gly Ser Gln Glu Asp Trp
20 25 30

Pro Arg Glu Pro 35

<210> 138

<211> 264

<212> PRT

<213> Homo sapien

<400> 138

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 1 5 10 15

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20 25 30

Ser Gly His Ala Ser Ser Thr Pro Gly Glu Lys Glu Thr Ser Ala 35 40 45

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Phe Asn 50 55 60

Ser Ser Leu Glu Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg 65 70 75 80

Asp Ile Ser Glu Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu 85 90 95

Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val Val Gln Leu 100 105 110

Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr

148

Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr

Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln 155

Ser Gly Ala Gly Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val 170 165

Cys Val Leu Val Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val 185 180

Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala 205 -200

Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His

Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys 235

Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala 250 245

Val Ala Ala Thr Ser Ala Asn Leu 260

<210> 139 <211> 241 <212> PRT <213> Homo sapien

<400> 139

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20 25

Ser Gly His Ala Ser Ser Thr Pro Gly Glu Glu Lys Glu Thr Ser Ala

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Phe Leu

149

Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe 70

Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly 85 90

Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr 100 105

Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser 115 120

Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val Pro Gly 130 135 140

Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala Leu Ala

Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg Lys Asn

Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met 180 185

Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser 195 200

Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly

Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn 230

Leu

<210> 140

<211> 92 <212> PRT

<213> Homo sapien

<400> 140

Met Ala Cys His Asn Leu Leu Ser Pro Gln Ala Val Cys Gln Cys Arg

Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr 20

150

His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val 35 40 45

Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Arg Leu Gly 50 55 60

Pro Thr Gly Gln Gly Lys Gln Arg Val Trp Leu Gly Lys Asp Ser Glu 65 70 75 80

Gly Gly Thr Trp Lys Thr Gln Arg Ala Trp Lys Arg 85 90

<210> 141

<211> 420

<212> PRT

<213> Homo sapien

<400> 141

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 1 10 15

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 20 25 30

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 35 40 45

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 50 55 60

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 65 70 75 80

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 85 90 95

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 100 105 110

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly
115 120 125

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 130 135 140

151

Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 145 150 155 160

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 165 170 175

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 180 185 190

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 195 200 205

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 210 215 220

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 225 230 235 240

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 245 250 255

Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe 260 265 270

Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln 275 280 285

Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val 290 295 300

Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly 305 310 315 320

Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val 325 330 335

Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr 355 360 365

His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val 370 375 380

Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly

152

400 395 390 385

Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr 410 405

Ser Ala Asn Leu 420

<210> 142

<211> 485 <212> PRT <213> Homo sapien

<400> 142

Met Pro Gln Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu

Thr Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr 20

Gly Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser

Ala Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val

Ser Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser

Ser Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro

Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro 105

Val Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val

Thr Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro 135

Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser 150 155 145

Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro 170 165

- Ala Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser 180 185 190
- Gly Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser 195 200 205
- Ala Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile 210 215 220
- Pro Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr 225 230 235 240
- Lys Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr 245 250 255
- Ser Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe 260 265 270
- Phe Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu 275 280 285
- Glu Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser 290 295 300
- Glu Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser 305 310 315
- Asn Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala 325 330 335
- Phe Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn 340 345 350
- Gln Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp 355 360 365
- Val Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala 370 375 380
- Gly Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu 385 390 395 400
- Val Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys 405 410 415

154

Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr 425 420

Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr

Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala 460 455

Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala 470

Thr Ser Ala Asn Leu 485

<210> 143 <211> 255 <212> PRT <213> Homo sapien

<400> 143

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 25 20

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser 35

Thr Glu Lys Asn Ala Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp 50

Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu Met Phe Leu Gln Ile

Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro 85

Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile 105 100

Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala

Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val 135 140

Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val Pro Gly Trp Gly 150

Ile Ala Leu Val Leu Val Cys Val Leu Val Ala Leu Ala Ile Val

Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly 185

Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu 200 195

Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr 215 210

Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser 225 230

Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu 250

<210> 144

<211> 517 <212> PRT <213> Homo sapien

<400> 144

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20 25

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 35

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85 90

156

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 180 185 190

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200 205

Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 275 280 285

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 290 295 300

Met Val Ser Ile Gly Leu Ser Phe Pro Ser Ser Pro Glu Ala Ala Ile 305 310 315 320

Arg Thr Val His Thr Leu Cys Ile Lys Pro Glu Ser Phe Pro Ser His 325 330 335

Pro Ser Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser

157

340 345 350

Asn Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala 355 360 365

Phe Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn 370 375 380

Gln Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp 385 390 395 400

Val Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala 405 410 415

Gly Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu 420 425 430

Val Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys 435 440 445

Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr 450 455 460

Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr 465 470 475 480

Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala 485 490 495

Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala 500 505 510

Thr Ser Ala Asn Leu 515

<210> 145

<211> 180

<212> PRT

<213> Homo sapien

<400> 145

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 1 5 10 15

Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe 20 25 30

158

Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln 40

Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val

Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly 75

Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val 85

Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg

Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr 120

His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val

Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly 155 150

Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr 165

Ser Ala Asn Leu

<210> 146 <211> 232 <212> PRT <213> Homo sapien

<400> 146

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Leu Thr 5

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 25 20

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser 35

Thr Glu Lys Asn Ala Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu

159

50 55 60

Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu 65 70 75 80

Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr 85 90 95

Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr 100 105 110

Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln 115 120 125

Ser Gly Ala Gly Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val 130 135 140

Cys Val Leu Val Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val 145 150 155 160

Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala 165 170 175

Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His 180 185 190

Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys
195 200 205

Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala 210 215 220

Val Ala Ala Thr Ser Ala Asn Leu 225 230

<210> 147

<211> 396

<212> PRT

<213> Homo sapien

<400> 147

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Leu Thr 1 5 10 15

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20 25 : 30

160

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 35 40 45

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50 55 60

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65 70 75 80

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85 90 95

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 180 185 190

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200 205

Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

161

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 280

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 295

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 315

Ile Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe 330 325

Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln

Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val 360

Ser Gly Glu Ala Thr Ser Leu Ala Ala Gln His His Ala Gly Ala 370

Pro Leu Leu Pro Val Ser Gly Ser Pro Leu Phe Pro 390

<210> 148 <211> 325 <212> PRT <213> Homo sapien

<400> 148

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 10 5

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20

Ser Gly His Ala Ser Ser Thr Pro Gly Glu Lys Glu Thr Ser Ala

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 55

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 90

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 100 105 110

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 180 185 190

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200 205

Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 275 280 285

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 290 295 300

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 305 310 315 320

Ile Lys Phe Ser Glu

<210> 149 <211> 409 <212> PRT <213> Homo sapien

<400> 149

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Leu Thr 5 10

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 25 20

Ser Gly His Ala Ser Ser Thr Pro Gly Glu Lys Glu Thr Ser Ala 35

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val 100

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 155 150

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 170

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 185 180

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 200 195

Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 

Ile Lys Phe Arg Pro Gly Ser Val Val Gln Leu Thr Leu Ala Phe 

Arg Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln 

Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val 

Ser Gly Cys Leu Ser Val Pro Pro Lys Glu Leu Arg Ala Ala Gly His 

Leu Ser Ser Pro Gly Tyr Leu Pro Ser Tyr Glu Arg Val Pro His Leu 

Pro His Pro Trp Ala Leu Cys Ala Pro 

<210> 150

<211> 379 <212> PRT <213> Homo sapien

<400> 150

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly
20 25 30

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala 35 40 45

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50 55 60

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65 70 75 80

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85 90 95

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val
100 105 110

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly
180 185 190

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200 205

Arg Ala Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

166

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 275 280 285

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 290 295 300

Met Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile 305 310 315 320

Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr 325 330 335

His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro 340 345 350

Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr 355 360 365

Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu 370 375

<210> 151

<211> 110

<212> PRT

<213> Homo sapien

<400> 151

Val Val Thr Trp His Asn Pro Gly Ala Gly Val Pro Gly Trp Gly Ile 1 5 10 15

Ala Leu Leu Val Leu Val Cys Val Leu Val Ala Leu Ala Ile Val Tyr
20 25 30

Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln 35 40 45

Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu Tyr 50 55 60

Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp 65 70 75 80

Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser Leu 85 90 95

167

Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu 105

<210> 152

<211> 127 <212> PRT <213> Homo sapien

<400> 152

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 25

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser

Thr Glu Lys Asn Ala Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly

Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu 70 75

Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr 85 90

Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser 100 105

Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Thr Ser Ala Asn Leu

<210> 153

<211> 336

<212> PRT <213> Homo sapien

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 5

Val Leu Thr Ala Thr Thr Ala Pro Lys Pro Ala Thr Val Val Thr Gly 20 25

Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser Ala

45

168

35

Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Val Ser 50 55 60

Met Thr Ser Ser Val Leu Ser Ser His Ser Pro Gly Ser Gly Ser Ser 65 70 75 80

40

Thr Thr Gln Gly Gln Asp Val Thr Leu Ala Pro Ala Thr Glu Pro Ala 85 90 95

Ser Gly Ser Ala Ala Thr Trp Gly Gln Asp Val Thr Ser Val Pro Val  $\cdot$  100 105 110

Thr Arg Pro Ala Leu Gly Ser Thr Thr Pro Pro Ala His Asp Val Thr 115 120 125

Ser Ala Pro Asp Asn Lys Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala 130 135 140

His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr 145 150 155 160

Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn Arg Pro Ala 165 170 175

Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser Ala Ser Gly 180 185 190

Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly Thr Ser Ala 195 200 205

Arg Ala Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro 210 215 220

Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys 225 230 235 240

Thr Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser 245 250 255

Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 260 265 270

Phe Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu 275 280 285

Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu 295 300

Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn 315 310

Ile Lys Phe Ser Gln Glu Leu Trp Trp Gln Asn Lys Arg Ser Ser Asn 330 325

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<213> Homo sapien

<400> 154

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Pro His Asp Pro Ser Ser Leu Gly Gly Met His Pro Ser Ser Val Ser 20 25

His Phe Arg Ala Phe Cys Thr Leu Leu Thr Leu Ser Arg Ile Pro Ala 35

Ile Trp Val Gln Ala Ser Gln 50

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<213> Homo sapien

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Ala Val Leu Gly Met Leu Arg Val Gln Ala Glu Ala Arg Glu Ala Gly 20 30

Leu Arg Gly Gln Lys Gln Gly Leu Gly Gln Ala Ser Pro Asp Arg Glu 35

Glu Gly Leu Arg Ser Lys Pro Ala Val Leu Leu Ala Gly Gly Pro Gly 50 55

Gly Tyr Lys Leu Leu Cys Ala Leu His Lys Pro Glu Ser Pro Thr Arg

170

70 65 75

Glu Asp Val Cys Glu Glu Pro Glu Thr Leu Asn Pro Ile Lys Arg Glu 90

Lys

<210> 156

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<400> 156

Met Leu Cys Ala Ile Ser Ile Ser Leu Val Ile Phe Phe Asn Lys His 10

Glu Ser Ile Lys Lys Lys Arg Arg Lys Lys Ala Gly Gly Thr Leu

Gly Gln Ser Gly Pro Gly Gly Asp Trp Phe Pro Ala Asn Ser Thr Ile

23

23

Ser Arg Thr Arg

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atcctgaatt ctgagaccat cca

<210> 159

<211> 21

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| 171 |
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| •              | •                       |    |  |  |
|                | 160                     | ٥. |  |  |
| agccgg.        | agga gatgtggete taccg   | 25 |  |  |
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172

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| cccagce        | accaccacca a            |    |
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|                | 166                     |    |
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| J J            |                         |    |
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173

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